

Li_esli_analysis

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Data preprocessing

```
# find and list all the files starting with "L".
# Maybe add full regular expr for the files

filenames = list.files(pattern="^[L]")

myfiles = lapply(filenames, read_csv)

# add a variable for filename
myfiles <- Map(cbind, myfiles, from = filenames)

for (i in 1:length(myfiles)) {
  myfiles[[i]]$word_count <- as.numeric(myfiles[[i]]$word_count)
}

library(stringr)

# an empty object for the loop
data <- list()

# first, create the list variable in each list,
# then take all list to merge them in one

# subject_nr! in 27 need to change to 27
# subject_nr! in 38 need to change to 38

for (i in 1:length(filenames)) {
  myfiles[[i]]$subj_id <- str_sub(filenames[i], 1, 5)
  myfiles[[i]]$subject_nr <- str_sub(filenames[i], 2, 3)

  data <- bind_rows(data, myfiles[i])
}

# create a dataset
data <- data %>%
  select(subj_id,
         subject_nr,
         avg_rt,
         coding,
         condition,
         conditon,
         count_exp_sequence,
         main_asp,
         main_num,
         main_tense,
```

```

    num,
    response,
    response_time,
    response_time_next,
    sentence_rus,
    sub_asp,
    sub_num,
    sub_tense,
    sub_wo,
    subject_parity,
    time_exp_sequence,
    time_next,
    time_pause,
    time_response_1,
    word_count,
    from)

# there was a typo in the randomization table, that's why we have condition and
# conditon as variables in the dataset

data %>%
  mutate(condition = ifelse(is.na(conditon) == T, condition, conditon)) %>%
  select(-conditon) %>%
  mutate(word_count = ifelse(num == 42, 6, word_count)) -> data

# data is clean

# read file with demografic data

demogr <- read.csv('demographics.csv', sep = ';' ) %>%
  mutate(subj_id = as.factor(subj_id))

working_data <- data %>% mutate(subj_id = as.factor(subj_id)) %>%
  full_join(demogr, by = 'subj_id') %>% filter(exclude == "no", english == 1) %>%
  mutate(heritageness = droplevels(heritageness)) %>%
  mutate(
    subj_id = as.factor(subj_id),
    subject_nr = as.factor(subject_nr),
    coding = as.factor(coding),
    condition = as.factor(condition),
    main_asp = as.factor(main_asp),
    main_num = as.factor(main_num),
    main_tense = as.factor(main_tense),
    response = as.character(response),
    sub_asp = as.factor(sub_asp),
    sub_num = as.factor(sub_num),
    sub_tense = as.factor(sub_tense),
    sub_wo = as.factor(sub_wo),
    from = as.factor(from)
  ) %>%
  filter(coding != c("10distr", "2li", "2distr", #the training trials
                    "12esli", "12li"))

# descriptive stats

```

```

demogr %>%
  filter(exclude == "no", english == 1) %>%
  mutate(female = case_when(gender == "male" ~ 0,
                             gender == "female" ~ 1))%>%
  group_by(heritageness, age_group) %>%
  summarise(count_n = n(),
            mean_age=mean(age, na.rm = T),
            max_age=max(age, na.rm = T),
            min_age=min(age, na.rm = T),
            sd_age = sd(age, na.rm = T),
            female = sum(female, na.rm = T),
            mean_edu=mean(edu_years, na.rm = T),
            english = sum(english, na.rm = T),
            french = sum(french, na.rm = T),
            ukranian = sum(ukranian, na.rm = T),
            chinese = sum(chinese, na.rm = T),
            italian = sum(italian, na.rm = T),
            japanese = sum(japanese, na.rm = T),
            spanish = sum(spanish, na.rm = T),
            hebrew = sum(hebrew, na.rm = T),
            german = sum(german, na.rm = T),
            georgian = sum(georgian, na.rm = T)) -> demogr_summary

demogr_summary %>% select(heritageness, age_group, count_n, mean_age, min_age, max_age, female, mean_edu

```

```

## # A tibble: 4 x 8
## # Groups:   heritageness [2]
##   heritageness age_group count_n mean_age min_age max_age female mean_edu
##   <fct>         <fct>    <int>   <dbl>   <int>   <int>   <dbl>   <dbl>
## 1 no          old         5    50.8    42     74      3    16.3
## 2 no          young       15    22.3    18     31     12    15.6
## 3 yes         old        12    50.2    42     69      5    18.1
## 4 yes         young        5    24.4    20     29      1    13.9

```

```

demogr_summary %>% select(heritageness, age_group, count_n, english, french, ukranian, chinese, italian

```

```

## # A tibble: 4 x 13
## # Groups:   heritageness [2]
##   heritageness age_group count_n english french ukranian chinese italian
##   <fct>         <fct>    <int>   <int>   <int>   <int>   <int>   <int>
## 1 no          old         5      5      1      0      0      1
## 2 no          young       15     15      2      0      1      0
## 3 yes         old        12     12      4      2      1      0
## 4 yes         young        5      5      3      0      0      0
## # ... with 5 more variables: japanese <int>, spanish <int>, hebrew <int>,
## #   german <int>, georgian <int>

```

```

demogr %>%
  filter(exclude == "no") %>%
  filter(heritageness == "yes") %>%
  select(heritageness, age_group, Immigr_data_the_USA) %>%
  mutate(Immigr_data_the_USA = as.numeric(as.character(Immigr_data_the_USA))),

```

```

years_inUSA = 2019 - Immigr_data_the_USA) %>%
group_by(heritageness, age_group) %>%
summarise(count_n = n(),
           mean_y_InUSA = mean(years_inUSA),
           min_y_InUSA = min(years_inUSA),
           max_y_InUSA = max(years_inUSA))

```

```

## # A tibble: 2 x 6
## # Groups:   heritageness [1]
##   heritageness age_group count_n mean_y_InUSA min_y_InUSA max_y_InUSA
##   <fct>         <fct>     <int>     <dbl>      <dbl>      <dbl>
## 1 yes          old           12        22         10         39
## 2 yes          young           5        11.6        4         22

```

Analysis

We will first check the data without dropping the observations by RTs

Visualisation

!!! We threat the ordinal variable (response) as numeric for the sake of visualisation. Need to think

```

facet_labels <- c(
  `distr` = "Control",
  `esli` = "Esli",
  `li` = "Li"
)

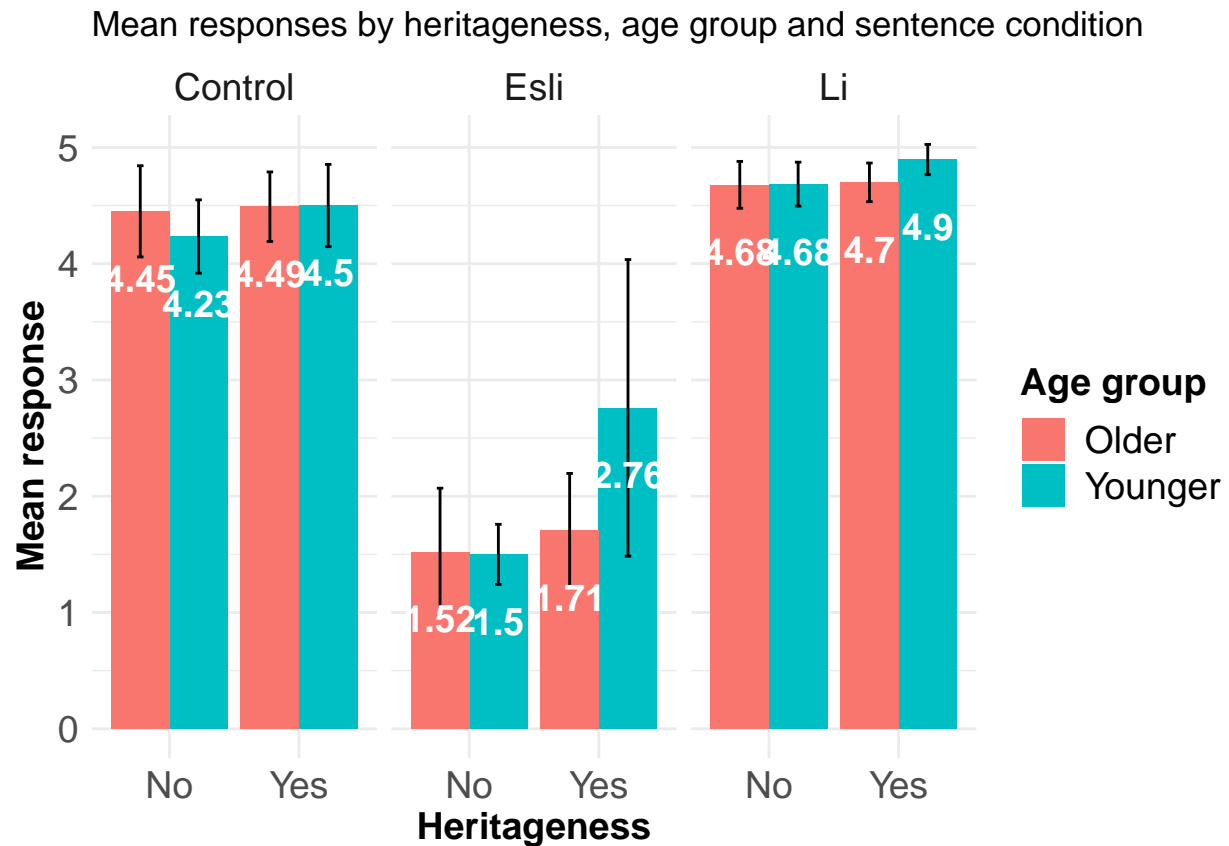
working_data %>% group_by(subj_id, condition, age_group, heritageness) %>%
  summarise(mean_subj = mean(as.numeric(response))) %>%
  group_by(condition, age_group, heritageness) %>%
  summarise(mean_resp = mean(mean_subj), sd_resp = sd(mean_subj), se_resp = sd_resp/sqrt(n())) %>%
  ggplot(aes(heritageness, mean_resp, fill = age_group))+
  geom_bar(stat = "identity", position=position_dodge())+
  geom_errorbar(aes(ymin = mean_resp - 2*se_resp,
                    ymax = mean_resp + 2*se_resp), width = 0.1, position=position_dodge(0.9))+
  geom_text(aes(label=round(mean_resp,2)), vjust=3, color="white",
            position = position_dodge(0.9), size=5, fontface = "bold" )+
#   scale_y_continuous(breaks = seq(0,1,0.05))+
  ggtitle("Mean responses by heritageness, age group and sentence condition")+
  facet_grid(.~condition, labeller = as_labeller(facet_labels))+
  theme_minimal()+
  scale_fill_discrete(labels = c("Older", "Younger"))+
  scale_x_discrete(labels = c("No", "Yes"))+
  theme(axis.text.x = element_text(size = 14),
        axis.text.y = element_text(size = 14),
        axis.title.x = element_text(size = 14, face = "bold"),
        axis.title.y = element_text(size = 14, face = "bold"),
        strip.text.x = element_text(size = 14),
        legend.title = element_text(size = 14, face = "bold"),

```

```

legend.text = element_text(size = 14)
) +
labs(x = "Heritageness", y = "Mean response", fill = "Age group")

```



Ordinal regression

```

# check the response variable
working_data$response[1:10]

```

```
## [1] "1" "5" "5" "2" "5" "5" "1" "4" "5" "1"
```

```
#now it is a factor variable, and we need an ordinal one
```

```

working_data$response_ordered <- ordered(working_data$response, levels = 1:5,
                                          labels = c('absolutely_unacceptable',
                                                    'mostly_unacceptable',
                                                    'indefinite',
                                                    'mostly_acceptable',
                                                    'absolutely_acceptable'))

```

```
working_data$response_ordered[1:10]
```

```
## [1] absolutely_unacceptable absolutely_acceptable
## [3] absolutely_acceptable mostly_unacceptable
## [5] absolutely_acceptable absolutely_acceptable
## [7] absolutely_unacceptable mostly_acceptable
## [9] absolutely_acceptable absolutely_unacceptable
## 5 Levels: absolutely_unacceptable < ... < absolutely_acceptable
```

```
test_data <- working_data %>% select(response_ordered,
                                     condition,
                                     age_group,
                                     subj_id,
                                     coding,
                                     heritageness) %>% arrange(subj_id)
```

```
# I first tried this packade but it took ages to compute
#install.packages("mixor")
#library(mixor)
#fit <- mixor(response ~ condition+age_group*heritageness,
#             data = test_data,
#             id = subj_id,
#             link = "logit")
#summary(fit)
```

```
#install.packages("ordinal")
library(ordinal)
```

```
## Warning: package 'ordinal' was built under R version 3.5.3
```

```
##
## Attaching package: 'ordinal'
```

```
## The following object is masked from 'package:dplyr':
```

```
##
## slice
```

```
mod <- clmm(response_ordered~condition*age_group*heritageness+(1|subj_id)+(1|coding), data=test_data,
            Hess=T
)
```

```
summary(mod)
```

```
## Cumulative Link Mixed Model fitted with the Laplace approximation
```

```
##
```

```
## formula: response_ordered ~ condition * age_group * heritageness + (1 |
## subj_id) + (1 | coding)
```

```
## data: test_data
```

```
##
## link threshold nobs logLik AIC niter max.grad cond.H
## logit flexible 1018 -916.12 1866.25 1614(4917) 1.54e-03 7.6e+02
##
## Random effects:
## Groups Name Variance Std.Dev.
## coding (Intercept) 0.3253 0.5703
## subj_id (Intercept) 0.4936 0.7026
## Number of groups: coding 47, subj_id 37
##
## Coefficients:
## Estimate Std. Error z value
## conditionesli -5.4718 0.5643 -9.696
## conditionli 0.9016 0.5759 1.566
## age_groupyoung -0.3917 0.5379 -0.728
## heritagenessyes 0.3168 0.5668 0.559
## conditionesli:age_groupyoung 0.6143 0.5420 1.133
## conditionli:age_groupyoung 0.2423 0.5953 0.407
## conditionesli:heritagenessyes 0.3802 0.5709 0.666
## conditionli:heritagenessyes -0.4828 0.6291 -0.767
## age_groupyoung:heritagenessyes 0.2293 0.7868 0.291
## conditionesli:age_groupyoung:heritagenessyes 1.4084 0.7656 1.839
## conditionli:age_groupyoung:heritagenessyes 0.7674 0.9175 0.836
## Pr(>|z|)
## conditionesli <2e-16 ***
## conditionli 0.1174
## age_groupyoung 0.4665
## heritagenessyes 0.5761
## conditionesli:age_groupyoung 0.2570
## conditionli:age_groupyoung 0.6840
## conditionesli:heritagenessyes 0.5054
## conditionli:heritagenessyes 0.4429
## age_groupyoung:heritagenessyes 0.7708
## conditionesli:age_groupyoung:heritagenessyes 0.0659 .
## conditionli:age_groupyoung:heritagenessyes 0.4029
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
## Estimate Std. Error z value
## absolutely_unacceptable|mostly_unacceptable -4.5253 0.5441 -8.318
## mostly_unacceptable|indefinite -2.9900 0.5274 -5.670
## indefinite|mostly_acceptable -2.0072 0.5177 -3.877
## mostly_acceptable|absolutely_acceptable -0.8393 0.5114 -1.641
```

Now we will try to cut off the responses with RTs which more then 20% bigger then median (within each participant)

```
# This is some trials to take into account the rt data
```

```
data_adjust_rt <- working_data %>%
  group_by(subj_id) %>%
  mutate(reference = median(response_time)*1.2) %>%
```

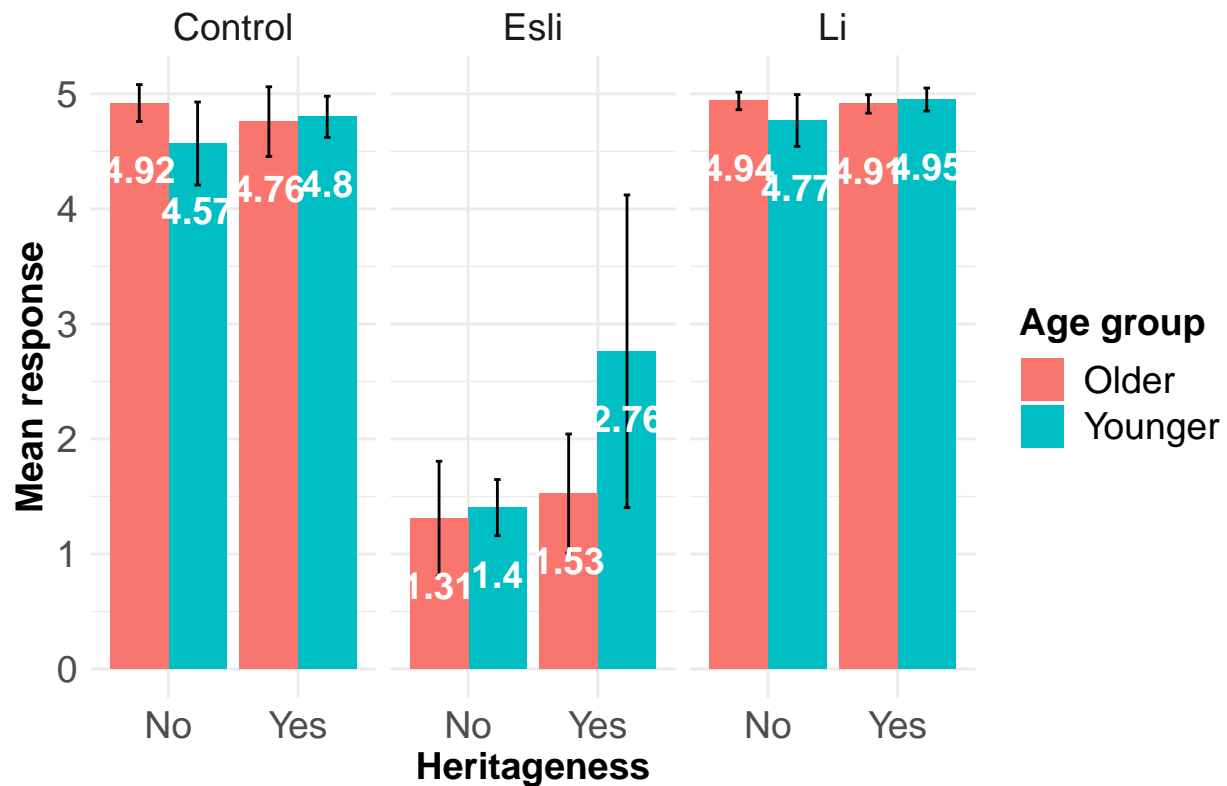
```

filter(response_time<=reference)

data_adjust_rt %>% group_by(subj_id, condition, age_group, heritageness) %>%
  summarise(mean_subj = mean(as.numeric(response))) %>%
  group_by(condition, age_group, heritageness) %>%
  summarise(mean_resp = mean(mean_subj), sd_resp = sd(mean_subj), se_resp = sd_resp/sqrt(n())) %>%
  ggplot(aes(heritageness, mean_resp, fill = age_group))+
  geom_bar(stat = "identity", position=position_dodge())+
  geom_errorbar(aes(ymin = mean_resp - 2*se_resp,
                    ymax = mean_resp + 2*se_resp), width = 0.1, position=position_dodge(0.9))+
  geom_text(aes(label=round(mean_resp,2)), vjust=3, color="white",
            position = position_dodge(0.9), size=5, fontface = "bold" )+
# scale_y_continuous(breaks = seq(0,1,0.05))+
  ggtitle("Mean responses by heritageness, age group and sentence condition (RT corrected data)")+
  facet_grid(.~condition, labeller = as_labeller(facet_labels))+
  theme_minimal()+
  scale_fill_discrete(labels = c("Older", "Younger"))+
  scale_x_discrete(labels = c("No", "Yes"))+
  theme(axis.text.x = element_text(size = 14),
        axis.text.y = element_text(size = 14),
        axis.title.x = element_text(size = 14, face = "bold"),
        axis.title.y = element_text(size = 14, face = "bold"),
        strip.text.x = element_text(size = 14),
        legend.title = element_text(size = 14, face = "bold"),
        legend.text = element_text(size = 14)
  ) +
  labs(x = "Heritageness", y = "Mean response", fill = "Age group")

```


Mean responses by heritageness, age group and sentence condition (RT cc)



```
# check the response variable
data_adjust_rt$response[1:10]
```

```
## [1] "5" "5" "5" "5" "1" "4" "1" "5" "1" "5"
```

```
#now it is a factor variable, and we need an ordinal one
```

```
data_adjust_rt$response_ordered <- ordered(data_adjust_rt$response, levels = 1:5,
                                           labels = c('absolutely_unacceptable',
                                                         'mostly_unacceptable',
                                                         'indefinite',
                                                         'mostly_acceptable',
                                                         'absolutely_acceptable'))
```

```
data_adjust_rt$response_ordered[1:10]
```

```
## [1] absolutely_acceptable absolutely_acceptable
## [3] absolutely_acceptable absolutely_acceptable
## [5] absolutely_unacceptable mostly_acceptable
## [7] absolutely_unacceptable absolutely_acceptable
## [9] absolutely_unacceptable absolutely_acceptable
## 5 Levels: absolutely_unacceptable < ... < absolutely_acceptable
```

```

test_data <- data_adjust_rt %>% select(response_ordered,
                                     condition,
                                     age_group,
                                     subj_id,
                                     coding,
                                     heritageness) %>%

  arrange(subj_id)

mod_rt <- clmm(response_ordered~condition*age_group*heritageness+
               (1|subj_id)+
               (1|coding), data=test_data,
               Hess=T
)

summary(mod_rt)

```

```

## Cumulative Link Mixed Model fitted with the Laplace approximation
##
## formula: response_ordered ~ condition * age_group * heritageness + (1 |
##      subj_id) + (1 | coding)
## data:      test_data
##
## link threshold nobs logLik AIC      niter      max.grad cond.H
## logit flexible  615  -385.18 804.36 1485(5870) 3.12e-04 1.2e+03
##
## Random effects:
## Groups Name      Variance Std.Dev.
## coding (Intercept) 7.302e-10 2.702e-05
## subj_id (Intercept) 1.489e+00 1.220e+00
## Number of groups: coding 47, subj_id 37
##
## Coefficients:
##                                     Estimate Std. Error z value
## conditionesli                      -8.4406      1.0335  -8.167
## conditionli                        0.2583      1.0918   0.237
## age_groupyoung                     -1.2005      1.0747  -1.117
## heritagenessyes                     -0.3673      1.1228  -0.327
## conditionesli:age_groupyoung        1.7419      1.0614   1.641
## conditionli:age_groupyoung          0.5797      1.2032   0.482
## conditionesli:heritagenessyes       1.4026      1.1059   1.268
## conditionli:heritagenessyes         0.1080      1.2749   0.085
## age_groupyoung:heritagenessyes      0.4951      1.4650   0.338
## conditionesli:age_groupyoung:heritagenessyes 1.6471      1.4036   1.173
## conditionli:age_groupyoung:heritagenessyes  0.2501      1.6658   0.150
##                                     Pr(>|z|)
## conditionesli                      3.16e-16 ***
## conditionli                        0.813
## age_groupyoung                     0.264
## heritagenessyes                     0.744
## conditionesli:age_groupyoung        0.101
## conditionli:age_groupyoung          0.630
## conditionesli:heritagenessyes       0.205

```

```
## conditionli:heritagenessyes 0.933
## age_groupyoung:heritagenessyes 0.735
## conditionesli:age_groupyoung:heritagenessyes 0.241
## conditionli:age_groupyoung:heritagenessyes 0.881
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
##
## Estimate Std. Error z value
## absolutely_unacceptable|mostly_unacceptable -6.7906 1.0290 -6.599
## mostly_unacceptable|indefinite -4.8370 0.9950 -4.861
## indefinite|mostly_acceptable -3.9944 0.9791 -4.080
## mostly_acceptable|absolutely_acceptable -2.8767 0.9618 -2.991
```

```
summary_mod <- summary(mod_rt)
```

```
data_summary <- data_adjust_rt %>% mutate(response=as.factor(response)) %>%
  group_by(subj_id, condition, age_group, heritageness) %>%
  mutate(all = n()) %>%
  group_by(subj_id, condition, age_group, heritageness, response, all) %>%
  summarise(quant = n()) %>%
  mutate(prop = quant/all) %>%
  ungroup() %>%
  group_by(condition, age_group, heritageness, response) %>%
  summarise(mean_prop = mean(prop), sd_prop = sd(prop), se_prop = sd_prop/sqrt(n()))
```

```
data_summary
```

```
## # A tibble: 42 x 7
## # Groups:   condition, age_group, heritageness [12]
##   condition age_group heritageness response mean_prop sd_prop se_prop
##   <fct>      <fct>      <fct>      <chr>      <dbl>    <dbl>    <dbl>
## 1 distr     old        no         4         0.4     NaN     NaN
## 2 distr     old        no         5         0.92    0.179    0.08
## 3 distr     old        yes        1         0.2     NaN     NaN
## 4 distr     old        yes        2         0.5     NaN     NaN
## 5 distr     old        yes        3         0.2     NaN     NaN
## 6 distr     old        yes        4         0.2     NaN     NaN
## 7 distr     old        yes        5         0.908    0.178    0.0514
## 8 distr     young      no         1         0.35    0.212    0.15
## 9 distr     young      no         2         0.25    NaN     NaN
## 10 distr    young      no         3         0.206    0.0419    0.0242
## # ... with 32 more rows
```

```
facet_labels <- c(
  `distr` = "Control",
  `esli` = "Esli",
  `li` = "Li",
  `yes` = "Heritage",
  `no` = "Non-heritage"
)
```

```
data_summary %>%
```

```

ggplot(aes(response, mean_prop, fill = age_group))+
  geom_bar(stat = "identity", position=position_dodge(0.9, preserve = "single"))+
  geom_errorbar(aes(ymin = mean_prop - 2*se_prop,
                    ymax = mean_prop + 2*se_prop), width = 0.13,
                position=position_dodge(0.9, preserve = "single"), color = "black")+
  facet_grid(heritageness~condition, labeller = as_labeller(facet_labels)) +
  geom_text(aes(label=round(mean_prop,2)), vjust=1, color="white",
            position = position_dodge(width = 0.9), size=2, fontface = "bold")+
  ggtitle("Mean response proportions by heritageness, age group and sentence condition")+
  theme_minimal()+
  theme(axis.text.x = element_text(size = 12),
        axis.text.y = element_text(size = 12),
        axis.title.x = element_text(size = 12, face = "bold"),
        axis.title.y = element_text(size = 12, face = "bold"),
        strip.text.x = element_text(size = 12),
        strip.text.y = element_text(size = 12),
        legend.title = element_text(size = 12),
        legend.text = element_text(size = 12),
        plot.title = element_text(size = 12)) +
  labs(x = "Response", y = "Mean proportion", fill = "Age group")+
  scale_fill_brewer(labels = c("Older", "Younger"), palette = "Paired")

```

Warning: Removed 12 rows containing missing values (geom_errorbar).

