

pilot1

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uncomment below to render to html

output:

bookdown::html_document2:

toc: true

toc__depth: 4

theme: cosmo

highlight: tango

uncomment below to render to pdf

output: bookdown::pdf_book: toc: true toc__depth: 4 highlight: tango

bibliography: [references/packages.bib, references/references.bib] biblio-style: apalike nocite: '@*' —

Results

```
# additional libraries
library("knitr")
library("janitor")
library("broom.mixed")
library("lme4")
library("emmeans")
library("tidyverse")
library("kableExtra")
```

```
## Warning in !is.null(rmarkdown::metadata$output) && rmarkdown::metadata$output
## %in% : 'length(x) = 2 > 1' in coercion to 'logical(1)'
```

```
library("modelr")
library("broom")
library("nlme")
library("meta")
library("metafor")
library(jtools) # Load jtools
theme_set(theme_classic())
```

```
# reading the data file
pilot1_data = read_csv("252.csv")
df_shape= filter(pilot1_data, !is.na(d))
# pilot1_data = pilot1_data %>%
#   select(ID, Title, d, d_var, Author)

df_shape_summary = df_shape %>%
  group_by(ID, Title, Author) %>%
  summarize(mean = mean(d),
            mean_se = mean(d_var))
```

First, Visualizing the data to have an initial idea of how it looks. First dividing the language group into two groups: the first one is the indo-european group which includes the English and the Spanish languages. The second group includes the rest of the languages: Japanese, Chinese, Tsimane.

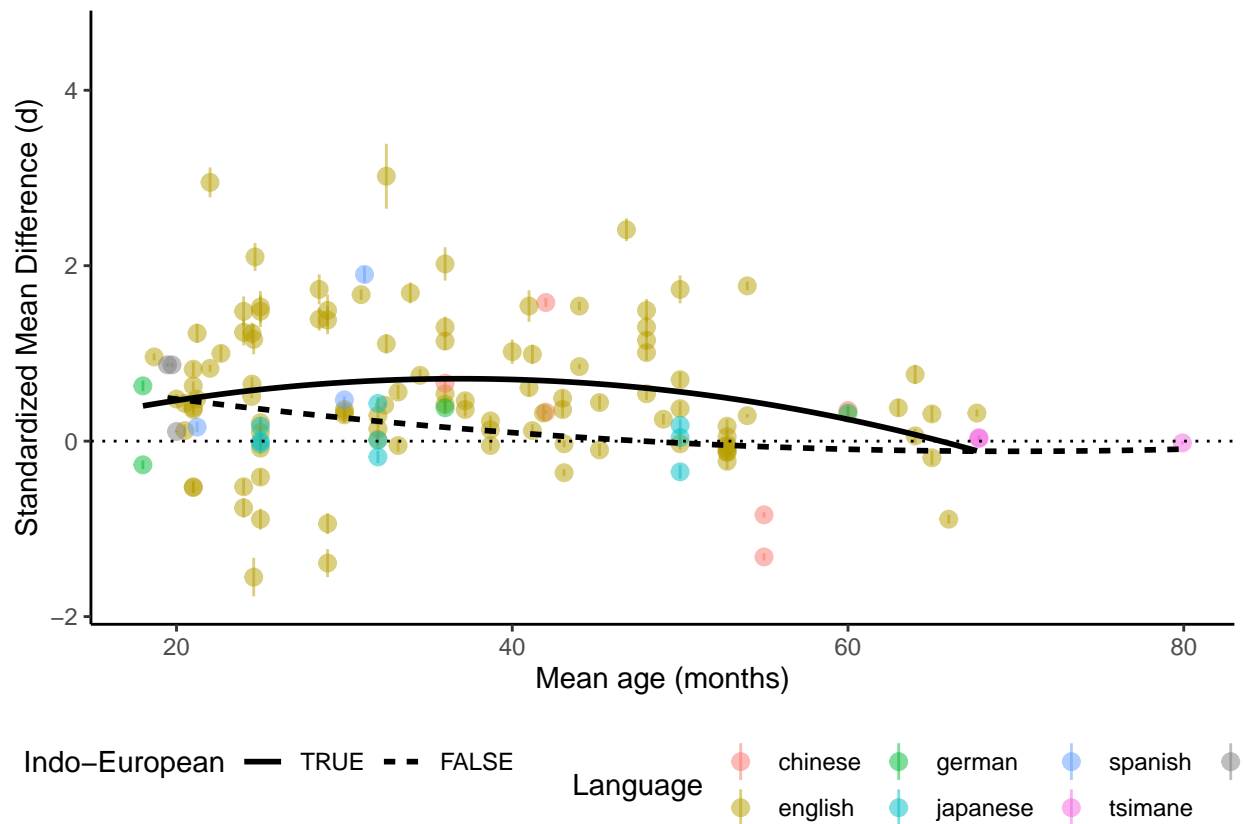
```
df_shape$indoeuropean <- fct_relevel(as.factor(df_shape$language %in%
                                              c("english","spanish", "german")),
                                     "TRUE")
```

creating a plot that shows the effects sizes colored per language group as well as the polynomial regression

```
ggplot(df_shape,
       aes(x = mean_age_months, y = d, color = language))+
  geom_pointrange(aes(ymin = d - d_var, ymax = d + d_var),
                 alpha = .5) +
  geom_smooth(aes(group = indoeuropean,
                  lty = indoeuropean),
              col = "black",
              method = "lm", se = FALSE,
              formula = y ~ poly(x,2)) +
  geom_hline(yintercept = 0, lty = 3) +
  ylab("Standardized Mean Difference (d)") +
  xlab("Mean age (months)") +
  scale_color_discrete(name = "Language") +
  scale_linetype_discrete(name = "Indo-European") +
  theme(legend.position = "bottom")
```

```
## Warning: Removed 8 rows containing non-finite values ('stat_smooth()').
```

```
## Warning: Removed 8 rows containing missing values ('geom_pointrange()').
```



```
ggsave("first_graph.png", width = 7, height = 4)
```

```
## Warning: Removed 8 rows containing non-finite values ('stat_smooth()').
## Removed 8 rows containing missing values ('geom_pointrange()').
```

using the meta-analytic function meta-gen which calculates the weights for each effects and confidence interval, pooled effect size, the heterogeneity.

```
m.gen <- metagen( TE= d,
                  seTE = d_var,
                  studlab = ID,
                  data = df_shape,
                  sm = "SMD",
                  fixed = FALSE,
                  random = TRUE,
                  method.tau = "REML",
                  hakn = TRUE,
                  title = "pilot shape bias meta-analysis"
)

summary(m.gen) ['TE']
```

```
## $TE
```

```
## [1] 0.32 0.03 -0.02 0.04 0.33 -0.84 1.58 -1.32 1.54 1.77 0.85 0.29
## [13] 1.54 0.76 0.61 0.06 0.33 0.36 0.31 0.36 0.49 -0.19 0.30 1.02
## [25] 0.38 -0.05 0.36 0.12 -0.10 0.12 0.65 1.39 1.11 0.56 0.46 0.99
## [37] 0.44 0.43 1.23 1.73 3.02 0.05 -0.06 0.17 -0.04 -0.23 -0.09 -0.13
## [49] -0.12 -0.36 -0.03 0.32 2.41 -0.89 0.51 0.96 0.41 1.90 -0.52 0.38
## [61] -0.53 0.82 0.63 0.36 0.21 0.29 0.70 0.10 0.14 0.37 -0.08 0.02
## [73] -0.03 0.17 0.43 0.18 0.00 0.01 0.04 -0.03 -0.18 -0.35 1.48 -0.89
## [85] 1.53 -0.41 1.48 -0.52 1.24 -0.76 1.49 -0.94 1.38 -1.39 0.75 1.69
## [97] 2.10 0.87 0.87 0.11 0.63 -0.27 1.23 0.16 2.95 0.83 0.54 1.01
## [109] 1.14 1.15 2.02 1.30 1.30 1.49 0.42 0.54 1.73 0.25 0.13 0.23
## [121] -0.05 0.48 0.48 1.00 0.66 0.38 0.35 0.32 1.16 -1.55 1.67 0.47
## [133] 1.81 4.02 2.06 1.56 1.42 1.38 0.73 0.34
```

```
m.gen["TE.fixed"]
```

```
## $TE.fixed
## [1] 0.317156
```

```
m.gen["TE.random"]
```

```
## $TE.random
## [1] 0.5474713
```

```
m.gen["w.random"]
```

```
## $w.random
## [1] 1.536920 1.538575 1.539759 1.538575 1.539759 1.538575 1.534797 1.536920
## [9] 1.534797 1.534797 1.538575 1.539759 1.467455 1.512511 1.517331 1.525665
## [17] 1.521718 1.521718 1.521718 1.521718 1.521718 1.525665 1.521718 1.495546
## [25] 1.517331 1.529164 1.525665 1.529164 1.529164 1.529164 1.525665 1.501610
## [33] 1.512511 1.525665 1.525665 1.512511 1.525665 1.529164 1.507268 1.475031
## [41] 1.272342 1.525665 1.525665 1.525665 1.525665 1.521718 1.525665 1.525665
## [49] 1.525665 1.536920 1.536920 1.521718 1.501610 1.534797 1.539759 1.536920
## [57] 1.538575 1.521718 1.529164 1.534797 1.532210 1.525665 1.534797 1.532210
## [65] 1.529164 1.529164 1.521718 1.529164 1.529164 1.525665 1.529164 1.529164
## [73] 1.529164 1.529164 1.529164 1.529164 1.529164 1.529164 1.529164 1.529164
## [81] 1.529164 1.525665 1.467455 1.507268 1.467455 1.521718 1.475031 1.521718
## [89] 1.489088 1.512511 1.467455 1.507268 1.482245 1.482245 1.529164 1.507268
## [97] 1.482245 1.539759 1.539759 1.540471 1.532210 1.534797 1.512511 1.532210
## [105] 1.475031 1.536920 1.529164 1.521718 1.517331 1.517331 1.459530 1.507268
## [113] 1.507268 1.501610 1.529164 1.529164 1.482245 1.532210 1.532210 1.532210
## [121] 1.532210 1.532210 1.532210 1.521718 1.529164 1.529164 1.532210 1.529164
## [129] 1.475031 1.433790 1.532210 1.536920 1.482245 1.026745 1.451269 1.495546
## [137] 1.501610 1.507268 1.525665 1.529164
```

forest plot using the m-gen function object

```
forextobj <- forest.meta(m.gen,
  sortvar = TE,
  prediction = TRUE,
  print.tau2 = FALSE,
  leftlabs = c("Author", "g", "SE"))
```

0.3300	0.0900	+	0.33	[0.15; 0.5
0.3400	0.0700	+	0.34	[0.20; 0.4
0.3500	0.0600	+	0.35	[0.23; 0.4
0.3600	0.0900	+	0.36	[0.18; 0.5
0.3600	0.0900	+	0.36	[0.18; 0.5
0.3600	0.0800	+	0.36	[0.20; 0.5
0.3600	0.0600	+	0.36	[0.24; 0.4
0.3700	0.0800	+	0.37	[0.21; 0.5
0.3800	0.1000	+	0.38	[0.18; 0.5
0.3800	0.0500	+	0.38	[0.28; 0.4
0.3800	0.0700	+	0.38	[0.24; 0.5
0.4100	0.0300	+	0.41	[0.35; 0.4
0.4200	0.0700	+	0.42	[0.28; 0.5
0.4300	0.0700	+	0.43	[0.29; 0.5
0.4300	0.0700	+	0.43	[0.29; 0.5
0.4400	0.0800	+	0.44	[0.28; 0.6
0.4600	0.0800	+	0.46	[0.30; 0.6
0.4700	0.0400	+	0.47	[0.39; 0.5
0.4800	0.0600	+	0.48	[0.36; 0.6
0.4800	0.0600	+	0.48	[0.36; 0.6
0.4900	0.0900	+	0.49	[0.31; 0.6
0.5100	0.0200	+	0.51	[0.47; 0.5
0.5400	0.0700	+	0.54	[0.40; 0.6

forest plot from the rma model:

```
forest(mod)
forest_data <-tibble(yi = mod$yi,
                     se = sqrt(mod$vi),
                     slab = mod$slab)

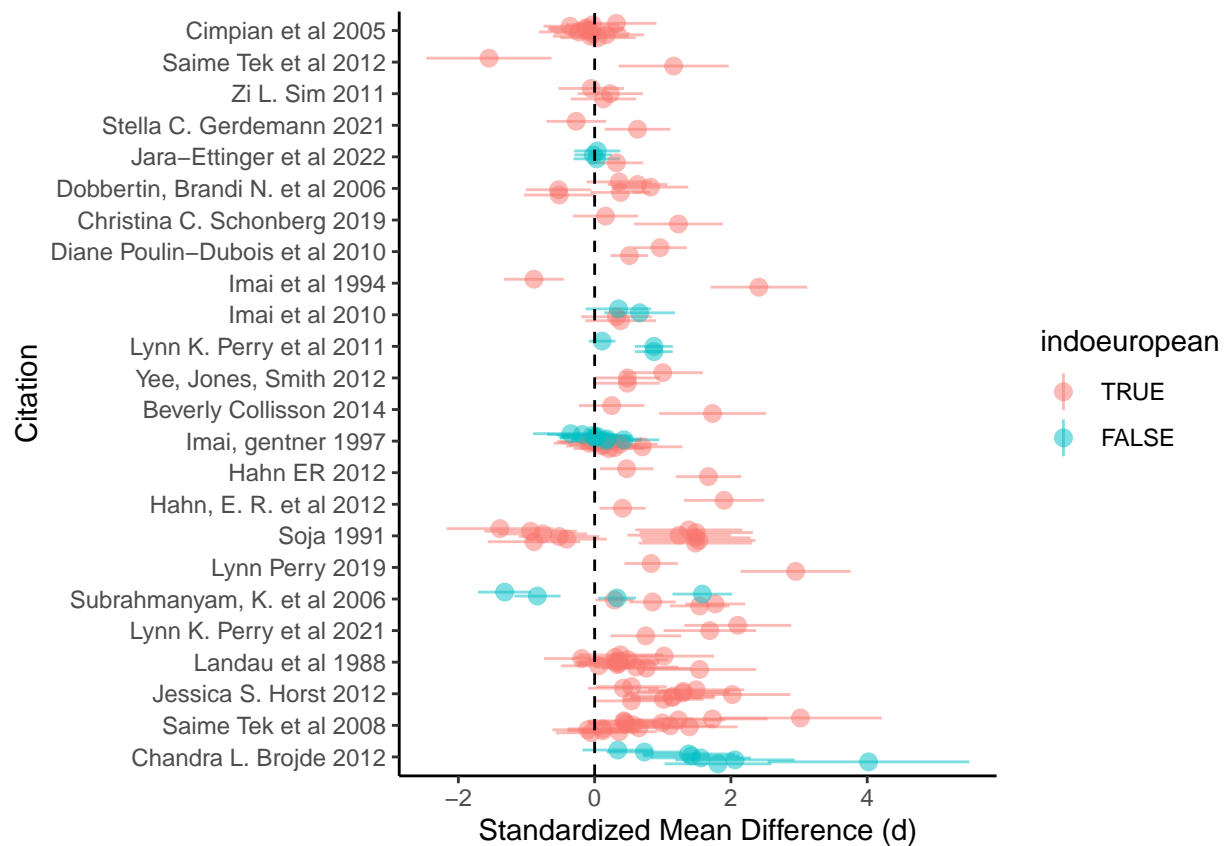
ggplot(forest_data,
       aes(x = slab, y = yi)) +
  geom_pointrange(aes(ymin = yi - se, ymax = yi + se)) +
  geom_hline(yintercept = 0, lty = 2) +
  coord_flip()

#
# theme_set(theme_bw(base_size=10))
# ata.frame(ES=ROM.ma$b, SE=ROM.ma$se, Type="Summary", Study="Summary"))
# forrest_data$Study2<-factor(forrest_data$Study, levels=rev(levels(forrest_data$Study)) )
# levels(forrest_data$Study2)
# plot1<-ggplot(data=forrest_data, aes(x=Study2, y=ES, ymax=ES+(1.96*SE), ymin=ES-(1.96*SE), size=factor(Typ
# plot2<-plot1+coord_flip()+geom_hline(aes(x=0), lty=2, size=1)+scale_size_manual(values=c(0.5,1))
# plot3<-plot2+xlab("Study")+ylab("log response ratio")+scale_colour_manual(values=c("grey", "black"))
# plot3+theme(legend.position="none")
```

#USING GG PLOT

```
df_shape$mean_age_months_centered36 <- df_shape$mean_age_months - 36
df_shape$log_mean_age_months <- log(df_shape$mean_age_months)

ggplot(df_shape, aes(x = short_cite, y = d,
                    ymin=d-sqrt(d_var)*1.96,
                    ymax=d+sqrt(d_var)*1.96)) +
  geom_pointrange(aes(color=indoeuropean), alpha = .5, position=position_dodge2(width=.5)) +
  coord_flip() +
  geom_hline(yintercept = 0, lty = 2) +
  aes(x=reorder(short_cite,-d, sum)) +
  ylab("Standardized Mean Difference (d)") +
  xlab("Citation")
```



```
#png("secondgraph.png")
```

Looking for Asymmetry

using funnel plots

```
col.contour = c("gray75", "gray85", "gray95")

funnel(m.gen,
```

```

comb.random = TRUE,
xlim = c(-2, 4),
contour = c(0.9, 0.95, 0.99),
col.contour = col.contour)

```

```

regtest(x = d, vi = d_var,
        data = df_shape)

```

```

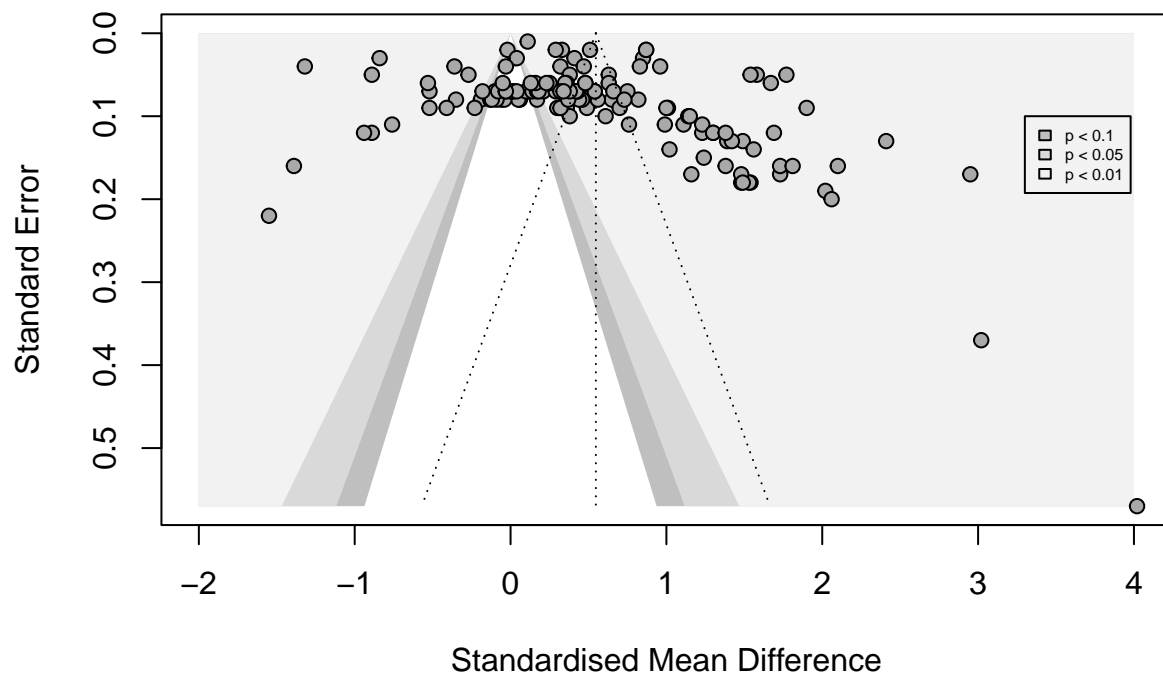
##
## Regression Test for Funnel Plot Asymmetry
##
## Model:      mixed-effects meta-regression model
## Predictor: standard error
##
## Test for Funnel Plot Asymmetry: z = 6.6402, p < .0001
## Limit Estimate (as sei -> 0):  b = -0.8597 (CI: -1.2814, -0.4381)

```

```

# Add a legend
legend(x = 3.3, y = 0.1, cex = 0.5,
       legend = c("p < 0.1", "p < 0.05", "p < 0.01"),
       fill = col.contour)

```



```

#png("funnel.png")

```

funnel plots using ggplot to account for moderators:

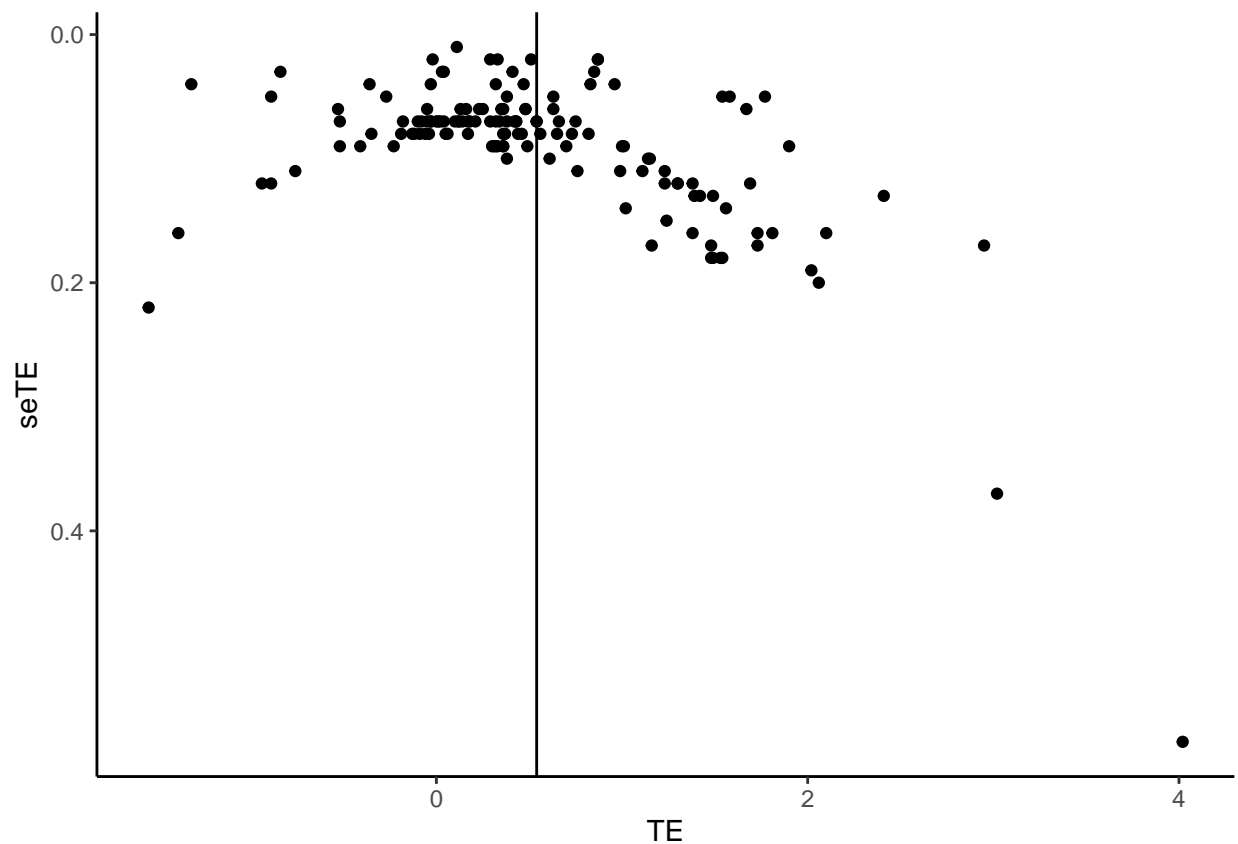
```
x = summary(m.gen)['TE']
y = summary(m.gen)['seTE']
m.gen["TE.fixed"]
```

```
## $TE.fixed
## [1] 0.317156
```

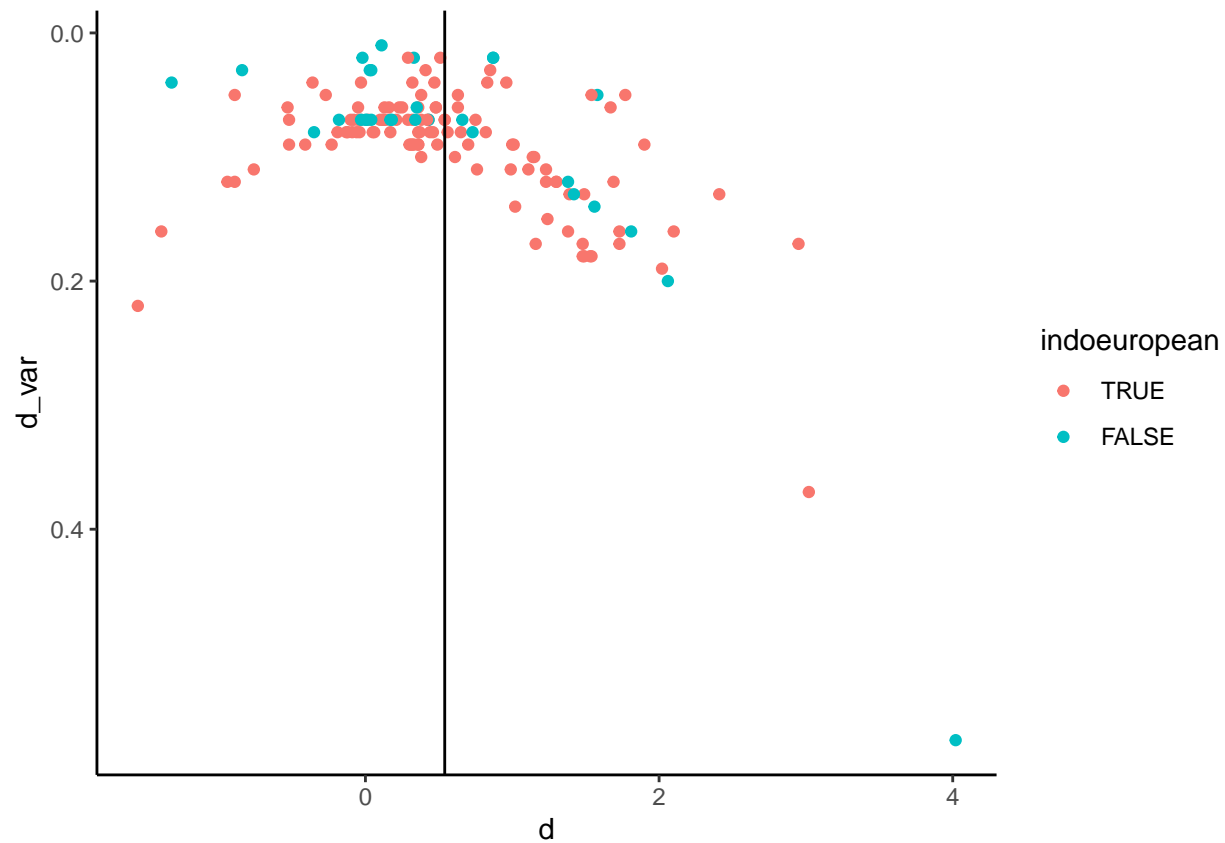
```
ter = m.gen["TE.random"]
```

```
data.gen = data.frame(x,y,ter)
```

```
ggplot(data = data.gen, mapping = aes(x=TE, y = seTE, color= )) +
  geom_point() +
  geom_vline(xintercept = 0.5401759) +
  scale_y_reverse()
```

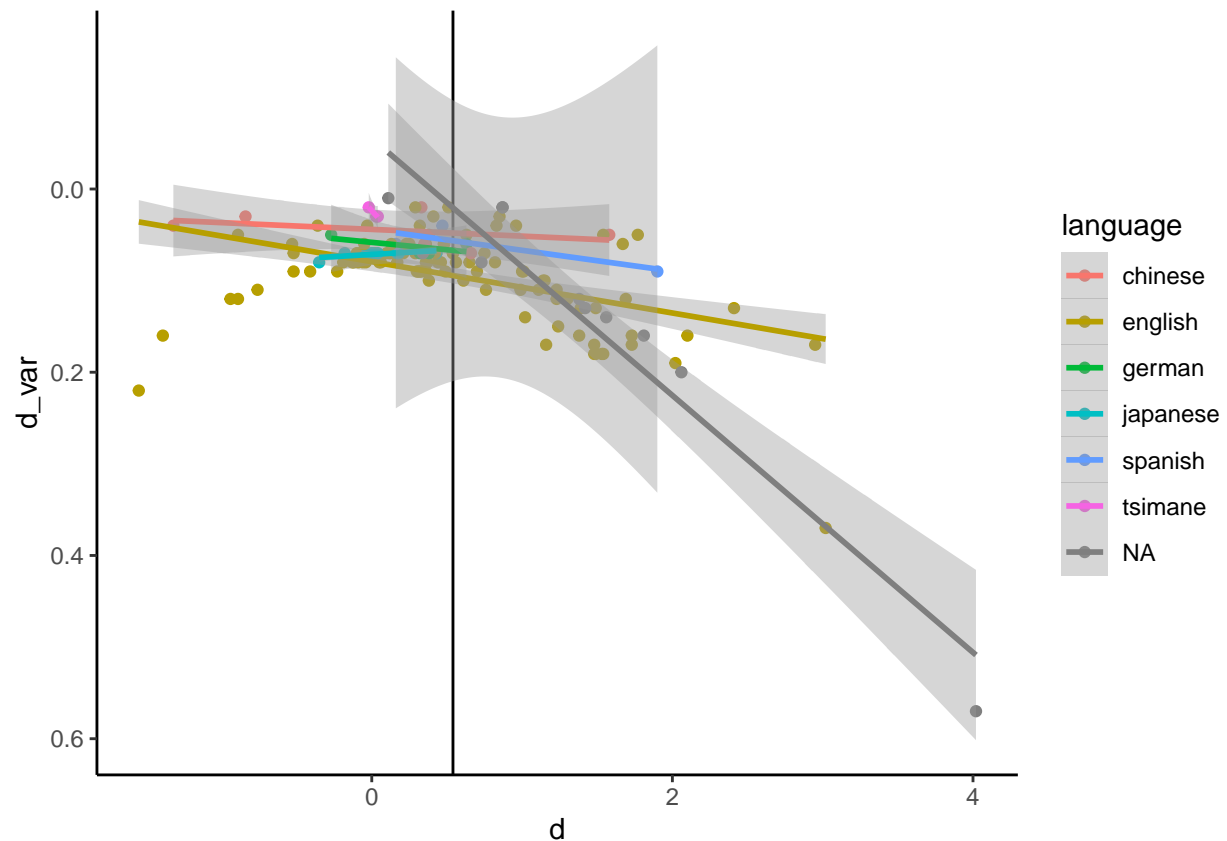


```
ggplot(data = df_shape, mapping = aes(x=d, y = d_var, color= indoeuropean)) +
  geom_point() +
  geom_vline(xintercept = 0.5401759) +
  scale_y_reverse()
```

```
ggplot(data = df_shape, mapping = aes(x=d, y = d_var, color= language)) +  
  geom_point() +  
  geom_vline(xintercept = 0.5401759) +  
  scale_y_reverse() +  
  geom_smooth(method = "lm")
```

```
## 'geom_smooth()' using formula = 'y ~ x'
```



```
# ggplot(data = df_shape, mapping = aes(x=d, y = d_var, color= agegroup)) +
#   geom_point() +
#   geom_vline(xintercept = 0.5401759) +
#   scale_y_reverse()

# +
#   aes(x=d, y=reorder(d_var, -d), color = indoeuropean)
```

Eggers regresstion test:

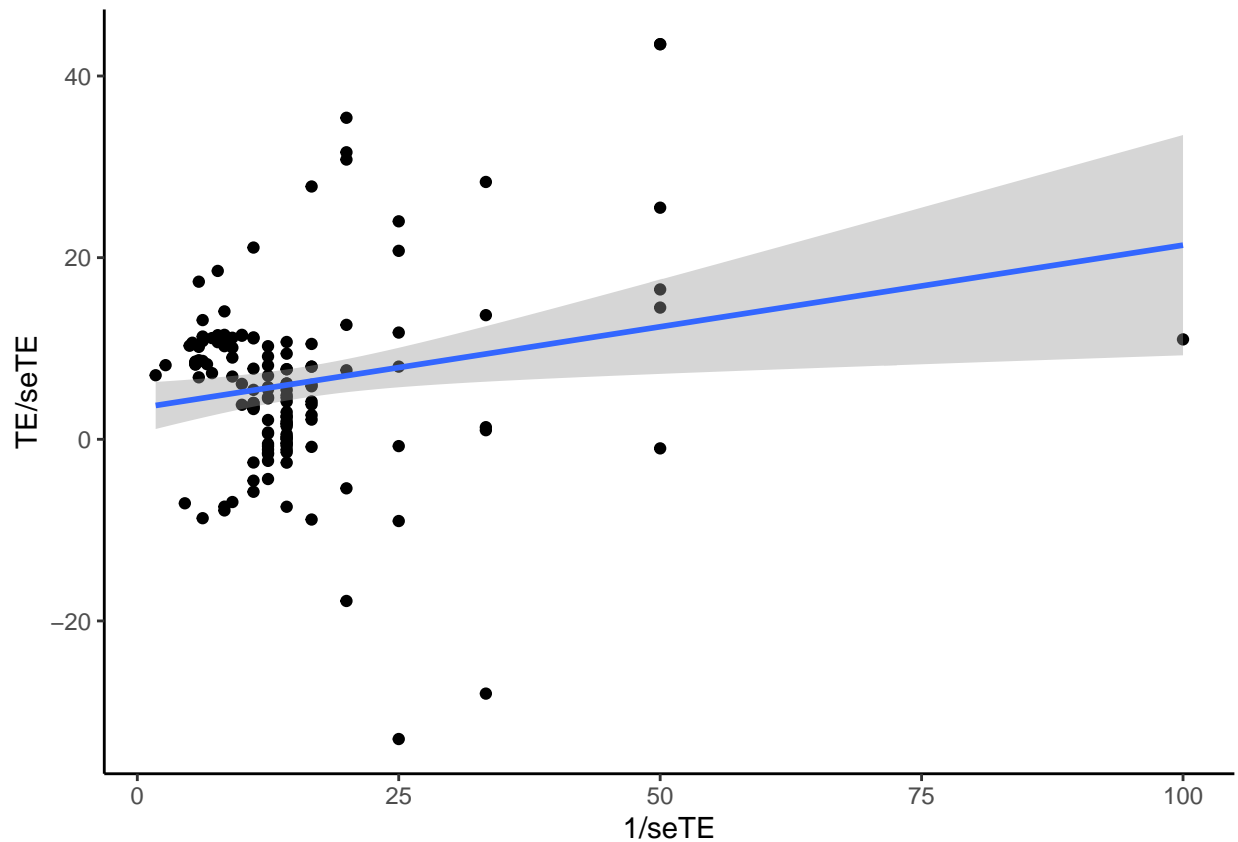
```
m.gen$data %>%
  mutate(y = m.gen$TE/m.gen$sseTE, x = 1/m.gen$sseTE) %>%
  lm(y ~ x, data= .) %>%
  summary()
```

```
##
## Call:
## lm(formula = y ~ x, data = .)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -40.900  -5.868   0.068   4.817  31.109
##
```

```
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)  3.40868    1.40767   2.422  0.0168 *
## x            0.17964    0.07184   2.500  0.0136 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 10.2 on 138 degrees of freedom
## Multiple R-squared:  0.04334,    Adjusted R-squared:  0.03641
## F-statistic: 6.252 on 1 and 138 DF,  p-value: 0.01357
```

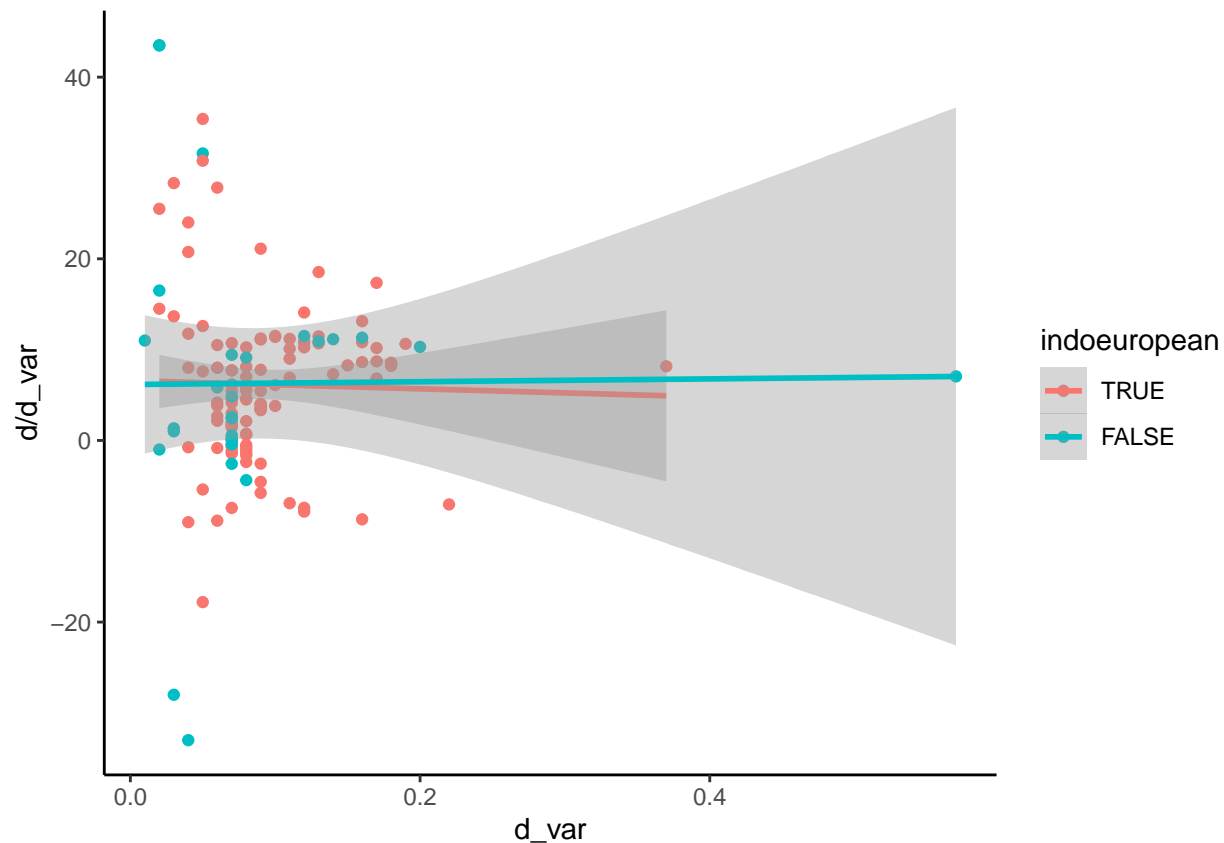
```
#eggerts regression
ggplot(data = data.gen, mapping = aes(x=1/seTE, y = TE/seTE, color= )) +
  geom_point() +
  geom_smooth(method = "lm")
```

```
## 'geom_smooth()' using formula = 'y ~ x'
```



```
ggplot(data = df_shape, mapping = aes(x=d_var, y = d/d_var, color= indoeuropean)) +
  geom_point() +
  geom_smooth(method = "lm")
```

```
## 'geom_smooth()' using formula = 'y ~ x'
```



using rma.mv instead of m.gen

```
mod <- rma.mv(yi = d,
              V = d_var,
              random = ~ 1 | ID,
              slab = short_cite,
              data = df_shape)

mod_nested <- rma.mv(yi = d,
                    V = d_var,
                    random = ~ 1 | ID/exp_num,
                    slab = short_cite,
                    data = filter(df_shape, !is.na(exp_num)))

summary(mod)
```

```
##
## Multivariate Meta-Analysis Model (k = 140; method: REML)
##
##      logLik   Deviance      AIC      BIC      AICc
## -345.7749   691.5499   695.5499   701.4188   695.6381
##
## Variance Components:
##
##      estim  sqrt  nlvls  fixed  factor
## sigma^2   0.1518 0.3897   25    no     ID
```

```
##
## Test for Heterogeneity:
## Q(df = 139) = 959.8543, p-val < .0001
##
## Model Results:
##
## estimate      se      zval      pval      ci.lb      ci.ub
##    0.4669    0.0828    5.6377    <.0001    0.3046    0.6293    ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
summary(mod_nested)
```

```
##
## Multivariate Meta-Analysis Model (k = 133; method: REML)
##
##      logLik      Deviance      AIC      BIC      AICc
## -328.8672    657.7344    663.7344    672.3828    663.9219
##
## Variance Components:
##
##      estim      sqrt      nlvls      fixed      factor
## sigma^2.1  0.1446  0.3803      25      no      ID
## sigma^2.2  0.0093  0.0964      34      no      ID/exp_num
##
## Test for Heterogeneity:
## Q(df = 132) = 895.1881, p-val < .0001
##
## Model Results:
##
## estimate      se      zval      pval      ci.lb      ci.ub
##    0.4670    0.0838    5.5735    <.0001    0.3028    0.6312    ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

plotting coefficients from the rmv model: assuming that those coefficients correspond to effect sizes

Confirmatory analysis

For primary analyses, i will exclude effect sizes from clinical populations and multilingual populations.

I will investigate the hypotheses via multi-level meta-regressions using the metafor package.

In all models, I will include random effects that control for non-independence between effect sizes based on grouping by paper and grouping by experiment.

I will first fit: Shape bias ~ 1 Shape bias ~ age shape bias ~ log(age) shape bias ~ poly(age,2)

intercept:

```
# using the meta and metafor packages to analyze meta-analysis effect sizes
mod_intercept <- rma.mv(d ~ 1,
  V = d_var,
  random = ~1 | as.factor>Title) /
  as.factor(exp_num),
  slab = Title,
  data = filter(df_shape, !is.na(exp_num)))

summary(mod_intercept)
```

```
##
## Multivariate Meta-Analysis Model (k = 133; method: REML)
##
##      logLik    Deviance      AIC      BIC      AICc
## -328.4943    656.9886    662.9886    671.6370    663.1761
##
## Variance Components:
##
##      estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.1519  0.3897    26    no      as.factor>Title)
## sigma^2.2  0.0027  0.0524    34    no  as.factor>Title)/as.factor(exp_num)
##
## Test for Heterogeneity:
## Q(df = 132) = 895.1881, p-val < .0001
##
## Model Results:
##
## estimate      se    zval    pval    ci.lb    ci.ub
##  0.4776  0.0828  5.7691  <.0001  0.3154  0.6399  ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

age

```
mod_age <- rma.mv(d ~ mean_age_months_centered36,
  V = d_var,
  random = ~1 | as.factor>Title) /
  as.factor(exp_num),
  slab = Title,
  data = filter(df_shape, !is.na(exp_num)))
```

```
## Warning: Rows with NAs omitted from model fitting.
```

```
summary(mod_age)
```

```
##
## Multivariate Meta-Analysis Model (k = 132; method: REML)
##
##      logLik    Deviance      AIC      BIC      AICc
```

```
## -311.2327    622.4654    630.4654    641.9355    630.7854
##
## Variance Components:
##
##           estim      sqrt  nlvls  fixed                                factor
## sigma^2.1  0.1143  0.3380    25     no                                as.factor(Title)
## sigma^2.2  0.0078  0.0884    33     no  as.factor(Title)/as.factor(exp_num)
##
## Test for Residual Heterogeneity:
## QE(df = 130) = 841.0606, p-val < .0001
##
## Test of Moderators (coefficient 2):
## QM(df = 1) = 27.8175, p-val < .0001
##
## Model Results:
##
##              estimate      se      zval      pval      ci.lb      ci.ub
## intrcpt              0.4438  0.0753   5.8971 <.0001    0.2963    0.5913
## mean_age_months_centered36 -0.0146  0.0028  -5.2742 <.0001   -0.0201   -0.0092
##
## intrcpt                ***
## mean_age_months_centered36 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

log age

```
mod_log_age <- rma.mv(d ~ log_mean_age_months,
  V = d_var,
  random = ~1 | as.factor(Title) /
    as.factor(exp_num),
  slab = Title,
  data = filter(df_shape, !is.na(log_mean_age_months)))

summary(mod_log_age)
```

```
##
## Multivariate Meta-Analysis Model (k = 132; method: REML)
##
##      logLik  Deviance      AIC      BIC      AICc
## -315.7124   631.4248   639.4248   650.8949   639.7448
##
## Variance Components:
##
##           estim      sqrt  nlvls  fixed                                factor
## sigma^2.1  0.1183  0.3439    25     no                                as.factor(Title)
## sigma^2.2  0.0059  0.0767    33     no  as.factor(Title)/as.factor(exp_num)
##
## Test for Residual Heterogeneity:
## QE(df = 130) = 851.6190, p-val < .0001
##
```

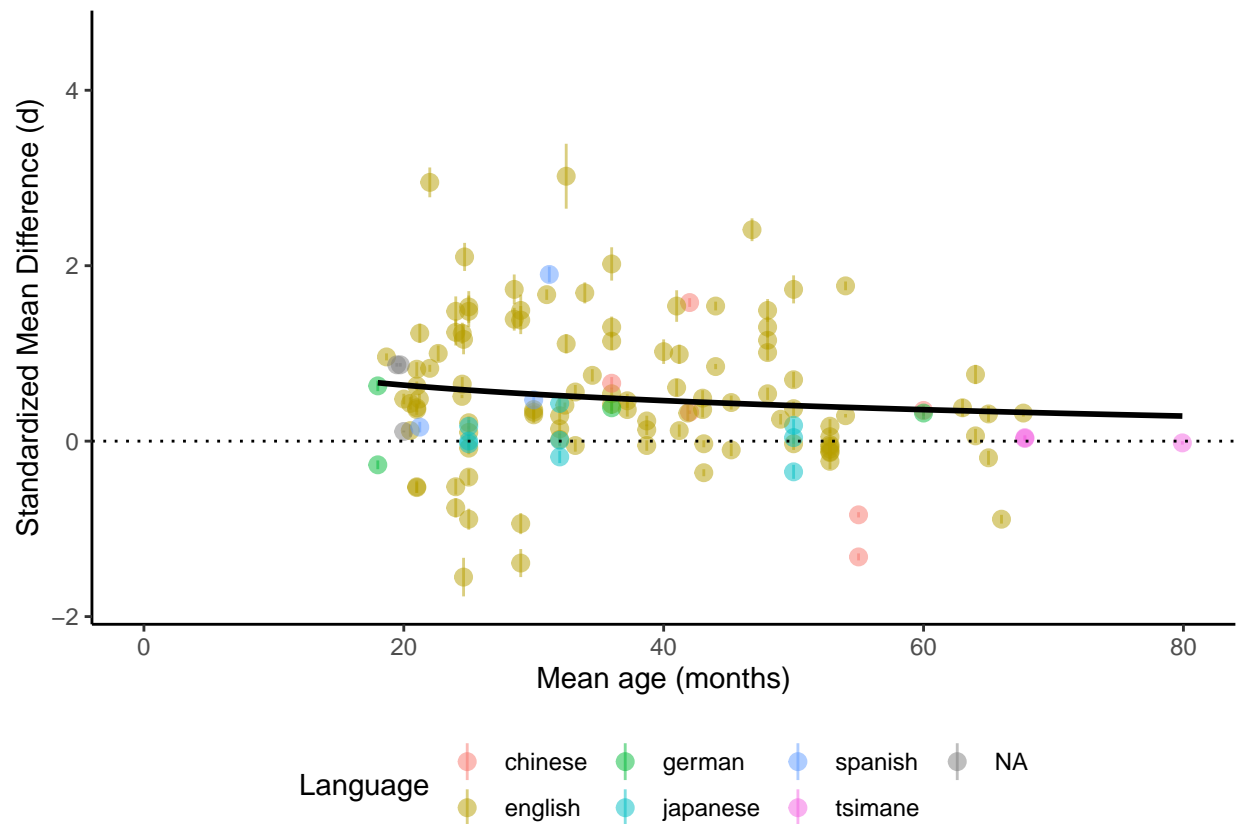
```
## Test of Moderators (coefficient 2):
## QM(df = 1) = 18.6968, p-val < .0001
##
## Model Results:
##
##               estimate      se      zval      pval      ci.lb      ci.ub
## intrcpt           2.0909  0.3879   5.3904 <.0001    1.3307    2.8512 ***
## log_mean_age_months -0.4711  0.1089  -4.3240 <.0001   -0.6846   -0.2575 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Let's look at what this means:

```
ggplot(df_shape,
       aes(x = mean_age_months, y = d, color = language)) +
  geom_pointrange(aes(ymin = d - d_var, ymax = d + d_var),
                 alpha = .5) +
  geom_smooth(aes(group = 1),
              col = "black",
              method = "lm", se = FALSE,
              formula = y ~ log(x)) +
  geom_hline(yintercept = 0, lty = 3) +
  ylab("Standardized Mean Difference (d)") +
  xlab("Mean age (months)") +
  scale_color_discrete(name = "Language") +
  scale_linetype_discrete(name = "Indo-European") +
  theme(legend.position = "bottom") +
  xlim(0,80)
```

```
## Warning: Removed 8 rows containing non-finite values ('stat_smooth()').
```

```
## Warning: Removed 8 rows containing missing values ('geom_pointrange()').
```

polynmoial age

```
mod_poly <- rma.mv(d ~ mean_age_months_centered36 + I(mean_age_months_centered36^2),
  V = d_var,
  random = ~1 | as.factor(Title) /
    as.factor(exp_num),
  slab = Title,
  data = filter(df_shape, !is.na(log_mean_age_months)))

summary(mod_poly)
```

```
##
## Multivariate Meta-Analysis Model (k = 132; method: REML)
##
##      logLik   Deviance      AIC      BIC      AICc
## -303.1769   606.3538   616.3538   630.6528   616.8416
##
## Variance Components:
##
##      estim  sqrt  nlvls  fixed      factor
## sigma^2.1  0.0984  0.3138    25    no  as.factor(Title)
## sigma^2.2  0.0423  0.2056    33    no as.factor(Title)/as.factor(exp_num)
##
## Test for Residual Heterogeneity:
```

```
## QE(df = 129) = 829.1803, p-val < .0001
##
## Test of Moderators (coefficients 2:3):
## QM(df = 2) = 45.6512, p-val < .0001
##
## Model Results:
##
##               estimate      se      zval      pval      ci.lb
## intrcpt           0.6197  0.0892   6.9485 <.0001    0.4449
## mean_age_months_centered36 -0.0048  0.0037  -1.2926  0.1961   -0.0121
## I(mean_age_months_centered36^2) -0.0007  0.0002  -4.1852 <.0001   -0.0011
##               ci.ub
## intrcpt           0.7945 ***
## mean_age_months_centered36    0.0025
## I(mean_age_months_centered36^2) -0.0004 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
m.gen.reg <- metareg(m.gen, ~language)
```

```
## Warning: Studies with NAs omitted from model fitting.
```

model comparison and plotting AICc , what is the criteria ? cutoff

```
#anova(mod_log_age, mod_poly, refit = TRUE)
#anova(mod_age, mod_poly) ## the two models are not comparable, not nested

#plot_component(mod, type = "BIC")
summary(mod_intercept)$fit.stats[5,'REML']
```

```
## [1] 663.1761
```

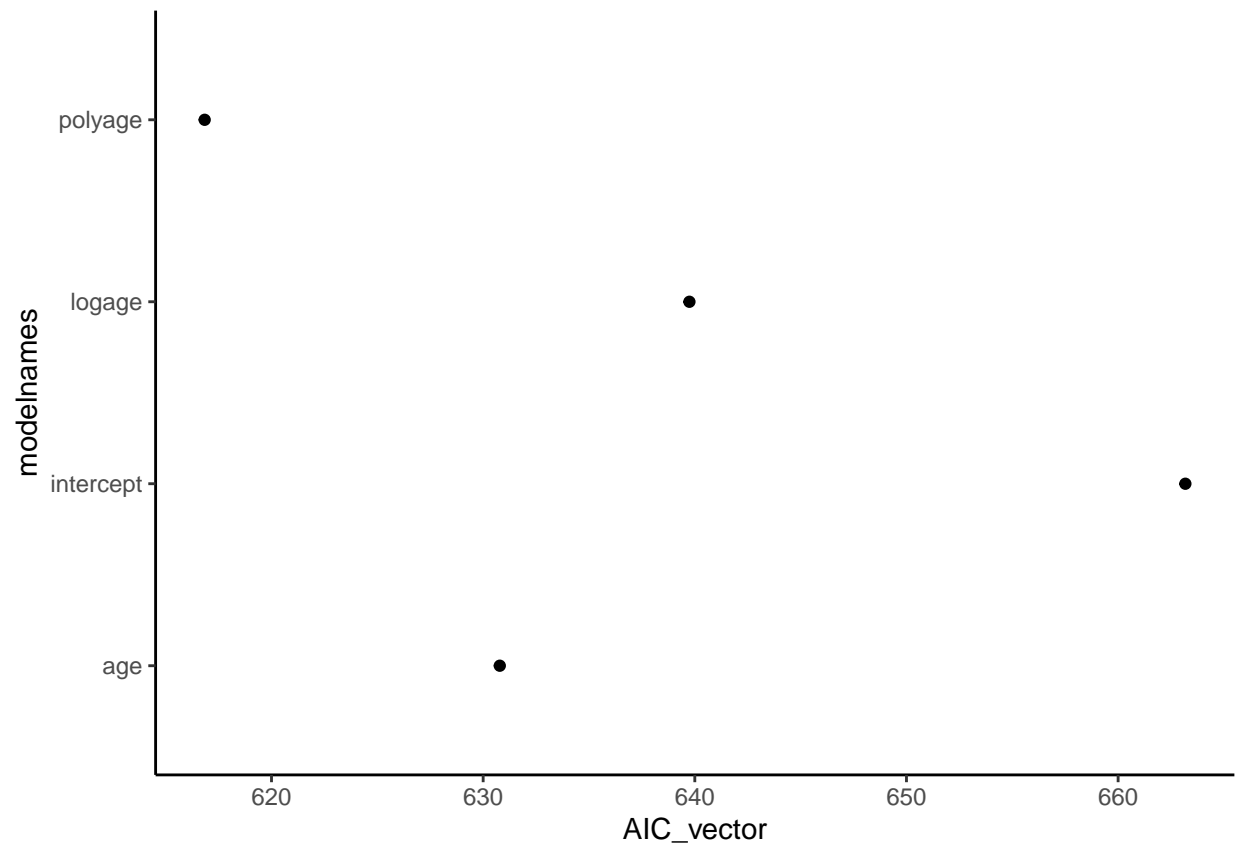
```
AIC_vector <- c(summary(mod_intercept)$fit.stats[5,'REML'],
                summary(mod_age)$fit.stats[5,'REML'],
                summary(mod_log_age)$fit.stats[5,'REML'],
                summary(mod_poly)$fit.stats[5,'REML'])

# AIC_vector <- c(AICc(mod_intercept), AICc(mod_age), AICc(mod_log_age), AICc(mod_poly))

modelnames <- c("intercept", "age", "logage", "polyage")

data.AIC = data.frame(AIC_vector, modelnames)

ggplot(data = data.AIC, mapping = aes(x = AIC_vector, y = modelnames)) +
  geom_point()
```



setting the contrasts:

```
#
# df_shape = df_shape %>%
#   mutate(lang_contrast = factor(language,
#                                   levels = c("english", "chinese", "german", "japanese",
#                                               "tsimane", "spanish"),
#                                   labels = c(0,-1,1,1,1,1)),
#   lang_contrast = lang_contrast %>%
#     as.character() %>%
#     as.factor() )
# contrasts(df_shape$lang_contrast) <- contr.sum(6)
# df_shape$lang_contrast

df_shape$lang_factor <- as.factor(df_shape$language)
# contrasts(df_shape$lang_factor) <- contr.treatment(6, base = 2)
df_shape$lang_factor <- fct_relevel(df_shape$lang_factor, "english", after = Inf)
contrasts(df_shape$lang_factor) <- contr.sum(6)*0.5
```

poly age with language

```
age_lang <- rma.mv(d ~ poly(mean_age_months_centered36,2) + language ,
                  V = d_var,
```

```

random = ~ 1 | ID/exp_num/language,
slab = short_cite,
data = filter(df_shape, !is.na(exp_num), !is.na(language)))

age_lang_interact <- rma.mv(d ~ poly(mean_age_months_centered36,2) * language,
V = d_var,
random = ~ 1 | ID/exp_num/language,
slab = short_cite,
data = filter(df_shape, !is.na(exp_num), !is.na(language)))

```

orthogonal polynomial:

Warning: Redundant predictors dropped from the model.

```
age_lang %>% summary()
```

```

##
## Multivariate Meta-Analysis Model (k = 129; method: REML)
##
##      logLik   Deviance      AIC      BIC      AICc
## -245.7844   491.5689   513.5689   544.3226   515.9909
##
## Variance Components:
##
##      estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0000  0.0001    23    no          ID
## sigma^2.2  0.0000  0.0000    32    no      ID/exp_num
## sigma^2.3  0.2284  0.4779    38    no ID/exp_num/language
##
## Test for Residual Heterogeneity:
## QE(df = 121) = 747.7508, p-val < .0001
##
## Test of Moderators (coefficients 2:8):
## QM(df = 7) = 60.5375, p-val < .0001
##
## Model Results:
##
##              estimate      se      zval      pval
## intrcpt              0.2658  0.3539   0.7509  0.4527
## poly(mean_age_months_centered36, 2)1 -2.4837  0.4782  -5.1942 <.0001
## poly(mean_age_months_centered36, 2)2 -2.4701  0.4897  -5.0445 <.0001
## languageenglish              0.2381  0.3675   0.6478  0.5171
## languagegerman              0.0743  0.5087   0.1460  0.8839
## languagejapanese            -0.3264  0.6015  -0.5426  0.5874
## languagespanish              0.3954  0.4747   0.8329  0.4049
## languagesimane              1.0230  0.4889   2.0925  0.0364
##              ci.lb      ci.ub
## intrcpt            -0.4279   0.9595
## poly(mean_age_months_centered36, 2)1 -3.4209 -1.5465 ***
## poly(mean_age_months_centered36, 2)2 -3.4298 -1.5103 ***
## languageenglish            -0.4823   0.9585
## languagegerman            -0.9228   1.0714
## languagejapanese          -1.5053   0.8525

```

```
## languagespanish          -0.5350    1.3258
## languagetsimane          0.0648    1.9813    *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
age_lang_interact %>% summary()
```

```
##
## Multivariate Meta-Analysis Model (k = 129; method: REML)
##
##      logLik    Deviance      AIC      BIC      AICc
## -225.6627    451.3255    491.3255    545.6955    500.5563
##
## Variance Components:
##
##      estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.1811  0.4255    23    no          ID
## sigma^2.2  0.0000  0.0000    32    no      ID/exp_num
## sigma^2.3  0.0511  0.2261    38    no ID/exp_num/language
##
## Test for Residual Heterogeneity:
## QE(df = 112) = 672.2426, p-val < .0001
##
## Test of Moderators (coefficients 2:17):
## QM(df = 16) = 110.5441, p-val < .0001
##
## Model Results:
##
##                                     estimate      se
## intrcpt                          0.4417    1.3571
## poly(mean_age_months_centered36, 2)1 -10.3159  11.4326
## poly(mean_age_months_centered36, 2)2  -2.5398  13.9288
## languageenglish                     0.1191   1.3708
## languagegerman                      -0.5372   1.1569
## languagejapanese                   -0.1032   1.4090
## languagespanish                    59.6411  16.7230
## languagetsimane                    -2.1232   7.1494
## poly(mean_age_months_centered36, 2)1:languageenglish  8.4474  11.4157
## poly(mean_age_months_centered36, 2)2:languageenglish  0.5605  13.9160
## poly(mean_age_months_centered36, 2)1:languagegerman   9.3529  12.3000
## poly(mean_age_months_centered36, 2)2:languagegerman   3.2744  15.0504
## poly(mean_age_months_centered36, 2)1:languagejapanese  9.1295  11.6640
## poly(mean_age_months_centered36, 2)2:languagejapanese  1.3671  14.4083
## poly(mean_age_months_centered36, 2)1:languagespanish  977.1301 271.1465
## poly(mean_age_months_centered36, 2)2:languagespanish  450.2977 129.0468
## poly(mean_age_months_centered36, 2)1:languagetsimane  20.1719  46.8774
##                                     zval    pval
## intrcpt                          0.3254  0.7449
## poly(mean_age_months_centered36, 2)1 -0.9023  0.3669
## poly(mean_age_months_centered36, 2)2 -0.1823  0.8553
## languageenglish                     0.0869  0.9308
## languagegerman                      -0.4643  0.6424
## languagejapanese                   -0.0732  0.9416
```

```
## languagespanish          3.5664  0.0004
## languagetsimane          -0.2970  0.7665
## poly(mean_age_months_centered36, 2)1:languageenglish  0.7400  0.4593
## poly(mean_age_months_centered36, 2)2:languageenglish  0.0403  0.9679
## poly(mean_age_months_centered36, 2)1:languagegerman    0.7604  0.4470
## poly(mean_age_months_centered36, 2)2:languagegerman    0.2176  0.8278
## poly(mean_age_months_centered36, 2)1:languagejapanese  0.7827  0.4338
## poly(mean_age_months_centered36, 2)2:languagejapanese  0.0949  0.9244
## poly(mean_age_months_centered36, 2)1:languagespanish    3.6037  0.0003
## poly(mean_age_months_centered36, 2)2:languagespanish    3.4894  0.0005
## poly(mean_age_months_centered36, 2)1:languagetsimane    0.4303  0.6670
##                               ci.lb      ci.ub
## intrcpt                               -2.2183      3.1016
## poly(mean_age_months_centered36, 2)1          -32.7234     12.0916
## poly(mean_age_months_centered36, 2)2          -29.8397     24.7601
## languageenglish                             -2.5676      2.8058
## languagegerman                             -2.8046      1.7303
## languagejapanese                           -2.8647      2.6584
## languagespanish                             26.8646     92.4176 ***
## languagetsimane                            -16.1357     11.8894
## poly(mean_age_months_centered36, 2)1:languageenglish -13.9269     30.8217
## poly(mean_age_months_centered36, 2)2:languageenglish -26.7144     27.8355
## poly(mean_age_months_centered36, 2)1:languagegerman -14.7547     33.4605
## poly(mean_age_months_centered36, 2)2:languagegerman -26.2238     32.7726
## poly(mean_age_months_centered36, 2)1:languagejapanese -13.7316     31.9905
## poly(mean_age_months_centered36, 2)2:languagejapanese -26.8727     29.6070
## poly(mean_age_months_centered36, 2)1:languagespanish  445.6927    1508.5675 ***
## poly(mean_age_months_centered36, 2)2:languagespanish  197.3706     703.2247 ***
## poly(mean_age_months_centered36, 2)1:languagetsimane -71.7061     112.0498
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

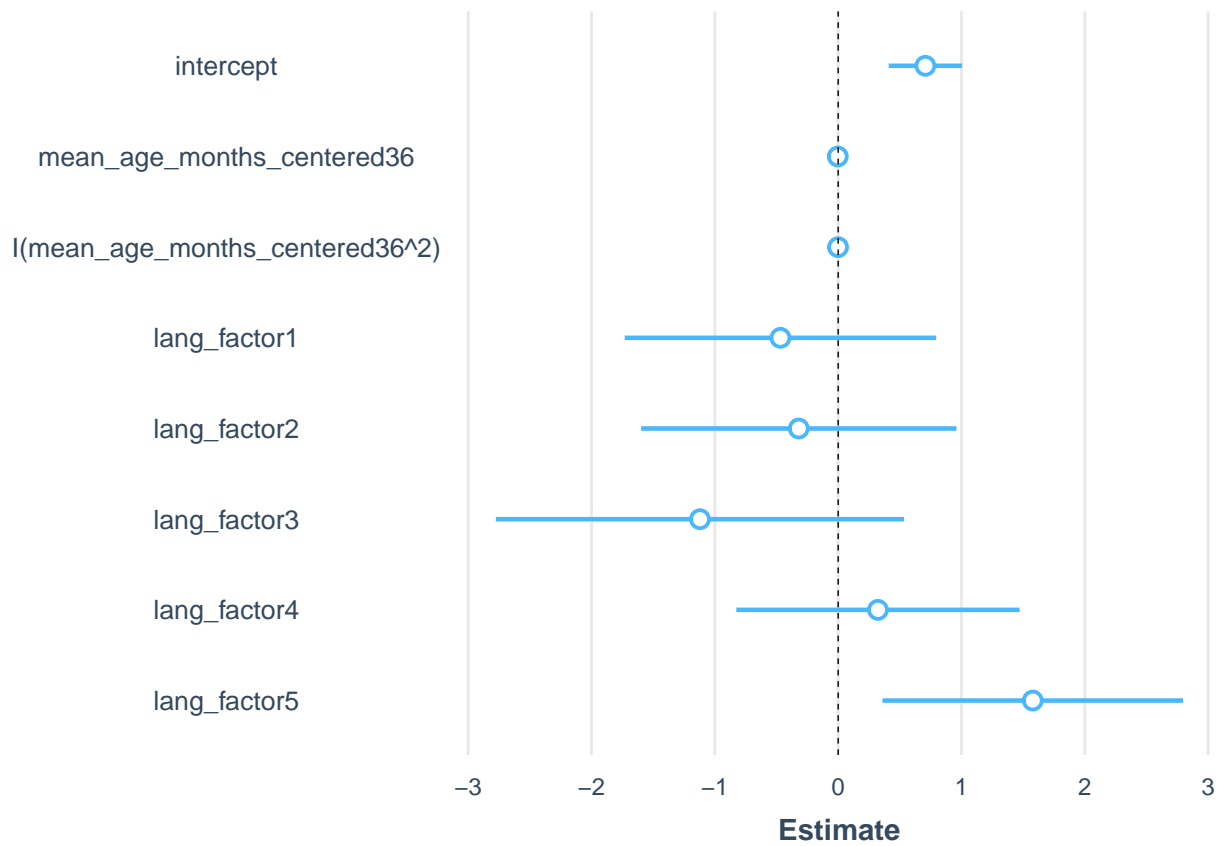
```
age_lang_npoly <- rma.mv(d ~ mean_age_months_centered36 + I(mean_age_months_centered36^2) + lang_factor,
  V = d_var,
  random = ~ 1 | ID/exp_num/language,
  slab = short_cite,
  data = filter(df_shape, !is.na(exp_num), !is.na(language)))

age_lang_interact_npoly <- rma.mv(d ~ mean_age_months_centered36 * lang_factor +
  I(mean_age_months_centered36^2) * lang_factor,
  V = d_var,
  random = ~ 1 | ID/exp_num/lang_factor,
  slab = short_cite,
  data = filter(df_shape, !is.na(exp_num), !is.na(language)))
```

the other polynomial:

```
## Warning: Redundant predictors dropped from the model.
```

```
plot_summs(age_lang_npoly, robust = TRUE)
```



```
plot_summs(age_lang_interact_npoly, robust = TRUE)
```



```
age_lang_npoly %>% summary()
```

```
##
## Multivariate Meta-Analysis Model (k = 129; method: REML)
##
##      logLik   Deviance      AIC      BIC      AICc
## -245.7844   491.5689   513.5689   544.3226   515.9909
##
## Variance Components:
##
##      estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.0000  0.0001   23    no          ID
## sigma^2.2  0.0000  0.0000   32    no      ID/exp_num
## sigma^2.3  0.2284  0.4779   38    no ID/exp_num/language
##
## Test for Residual Heterogeneity:
## QE(df = 121) = 747.7508, p-val < .0001
##
## Test of Moderators (coefficients 2:8):
## QM(df = 7) = 60.5375, p-val < .0001
##
## Model Results:
##
##      estimate      se      zval      pval      ci.lb
## intrcpt      0.7069  0.1518  4.6572 <.0001  0.4094
```



```

## mean_age_months_centered36      -0.0033  0.0040  -0.8159  0.4146  -0.0112
## I(mean_age_months_centered36^2) -0.0010  0.0002  -5.0445  <.0001  -0.0014
## lang_factor1                    -0.4681  0.6445  -0.7264  0.4676  -1.7313
## lang_factor2                    -0.3196  0.6522  -0.4900  0.6242  -1.5979
## lang_factor3                    -1.1209  0.8445  -1.3274  0.1844  -2.7761
## lang_factor4                     0.3226  0.5860   0.5506  0.5819  -0.8259
## lang_factor5                     1.5779  0.6218   2.5376  0.0112   0.3592
##                                ci.lb  ci.ub
## intrcpt                        1.0044  ***
## mean_age_months_centered36      0.0046
## I(mean_age_months_centered36^2) -0.0006  ***
## lang_factor1                     0.7951
## lang_factor2                     0.9588
## lang_factor3                     0.5342
## lang_factor4                     1.4712
## lang_factor5                     2.7967   *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```
age_lang_interact_npoly %>% summary()
```

```

##
## Multivariate Meta-Analysis Model (k = 129; method: REML)
##
##      logLik   Deviance      AIC      BIC      AICc
## -225.6627   451.3255   491.3255   545.6955   500.5563
##
## Variance Components:
##
##      estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.1811  0.4255    23    no          ID
## sigma^2.2  0.0000  0.0000    32    no      ID/exp_num
## sigma^2.3  0.0511  0.2261    38    no ID/exp_num/lang_factor
##
## Test for Residual Heterogeneity:
## QE(df = 112) = 672.2426, p-val < .0001
##
## Test of Moderators (coefficients 2:17):
## QM(df = 16) = 110.5441, p-val < .0001
##
## Model Results:
##
##      estimate      se      zval
## intrcpt          11.5861  3.2177  3.6007
## mean_age_months_centered36      0.1694  0.0584  2.9008
## lang_factor1     -21.6791  6.4096 -3.3823
## lang_factor2     -23.4460  6.4433 -3.6388
## lang_factor3     -22.2984  6.4700 -3.4465
## lang_factor4       9.6867  2.4130  4.0144
## lang_factor5      79.4585 23.6312  3.3624
## I(mean_age_months_centered36^2)    0.0368  0.0107  3.4326
## mean_age_months_centered36:lang_factor1 -0.4441  0.2321 -1.9137
## mean_age_months_centered36:lang_factor2 -0.3584  0.1317 -2.7207

```

```
## mean_age_months_centered36:lang_factor3      -0.3419   0.1172  -2.9172
## mean_age_months_centered36:lang_factor4       7.4139   2.0579   3.6026
## mean_age_months_centered36:lang_factor5      -5.9271   1.7075  -3.4711
## lang_factor1:I(mean_age_months_centered36^2)  -0.0758   0.0234  -3.2456
## lang_factor2:I(mean_age_months_centered36^2)  -0.0731   0.0216  -3.3801
## lang_factor3:I(mean_age_months_centered36^2)  -0.0747   0.0216  -3.4570
## lang_factor4:I(mean_age_months_centered36^2)   0.2989   0.0854   3.4989
##                                     pval      ci.lb      ci.ub
## intrcpt                                0.0003    5.2795   17.8926 ***
## mean_age_months_centered36             0.0037    0.0549    0.2838 **
## lang_factor1                           0.0007   -34.2416  -9.1166 ***
## lang_factor2                           0.0003   -36.0748 -10.8173 ***
## lang_factor3                           0.0006   -34.9792  -9.6175 ***
## lang_factor4                           <.0001    4.9573   14.4161 ***
## lang_factor5                           0.0008   33.1422  125.7749 ***
## I(mean_age_months_centered36^2)         0.0006    0.0158    0.0579 ***
## mean_age_months_centered36:lang_factor1     0.0557   -0.8989    0.0107 .
## mean_age_months_centered36:lang_factor2     0.0065   -0.6165   -0.1002 **
## mean_age_months_centered36:lang_factor3     0.0035   -0.5716   -0.1122 **
## mean_age_months_centered36:lang_factor4     0.0003    3.3804   11.4473 ***
## mean_age_months_centered36:lang_factor5     0.0005   -9.2738   -2.5804 ***
## lang_factor1:I(mean_age_months_centered36^2) 0.0012   -0.1216   -0.0300 **
## lang_factor2:I(mean_age_months_centered36^2) 0.0007   -0.1154   -0.0307 ***
## lang_factor3:I(mean_age_months_centered36^2) 0.0005   -0.1170   -0.0323 ***
## lang_factor4:I(mean_age_months_centered36^2) 0.0005    0.1314    0.4663 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

poly age with syntax

```
age_lang_syntax <- rma.mv(d ~ poly(mean_age_months,2) + language + Informative_syntax,
  V = d_var,
  random = ~ 1 | ID/exp_num/language,
  slab = short_cite,
  data = filter(df_shape, !is.na(exp_num), !is.na(language)))
```

```
## Warning: Rows with NAs omitted from model fitting.
```

```
rma.mv(d ~ poly(mean_age_months,2)*Informative_syntax,
  V = d_var,
  random = ~1 | as.factor(Title) +
    as.factor(exp_num),
  slab = Title,
  data = filter(df_shape, !is.na(mean_age_months))) %>% glance()
```

```
## Warning: Rows with NAs omitted from model fitting.
```

```
## Warning: One or more levels of inner factor (i.e., The shape-bias in
## Spanish-speaking children and its relationship to vocabulary) removed due to
## NAs.
```

```
## # A tibble: 1 x 12
##   tau.squ~1 cochr~2 p.valu~3 cochr~4 p.valu~5 df.re~6 logLik devia~7   AIC   BIC
##   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <int> <dbl>   <dbl> <dbl> <dbl>
## 1     0.214     801. 1.47e-99     53.1 3.24e-10     124 -294.    588.  604.  627.
## # ... with 2 more variables: AICc <dbl>, nobs <int>, and abbreviated variable
## #   names 1: tau.squared, 2: cochrane.qe, 3: p.value.cochrane.qe, 4: cochrane.qm,
## #   5: p.value.cochrane.qm, 6: df.residual, 7: deviance
```

Exploratory analysis

```
rma.mv(d ~ poly(mean_age_months,2),
      V = d_var,
      random = ~1 | as.factor(language)+
        as.factor>Title),
      slab = Title,
      data = filter(df_shape, !is.na(mean_age_months), !is.na(language))) %>% glance()
```

```
## # A tibble: 1 x 12
##   tau.squ~1 cochr~2 p.valu~3 cochr~4 p.valu~5 df.re~6 logLik devia~7   AIC   BIC
##   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <int> <dbl>   <dbl> <dbl> <dbl>
## 1     0.206     800. 1.59e-98     41.5 9.66e-10     126 -254.    508.  518.  532.
## # ... with 2 more variables: AICc <dbl>, nobs <int>, and abbreviated variable
## #   names 1: tau.squared, 2: cochrane.qe, 3: p.value.cochrane.qe, 4: cochrane.qm,
## #   5: p.value.cochrane.qm, 6: df.residual, 7: deviance
```

Discussion

References