

Shrinkage and partial pooling

Mixed Models 3

Daniela Palleschi

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Learning Objectives

Today we will learn...

- about no/complete/partial pooling
- about shrinkage

Resources

- this lecture covers
 - [Blog post](#) “Plotting partial pooling in mixed-effects models” from Tristin Mahr (2017)
 - Section 15.9 'Shrinkage and Individual Differences' in Winter (2019)
 - Box 8.2 'Broader Context: Shrinkage and Partial Pooling' in Sonderegger (2023)
- we will be using the data from Biondo et al. (2022)

Set-up

```
# suppress scientific notation
options(scipen=999)
```

Load packages

```
# load libraries
pacman::p_load(
  tidyverse,
  janitor,
  here,
  lmerTest)
```

```
lmer <- lmerTest::lmer
```

Load data

- data from Biondo et al. (2022)

```
df_biondo <-
  read_csv(here("data", "Biondo.Soilemezidi.Mancini_dataset_ET.csv"),
    locale = locale(encoding = "Latin1") ## for special characters in Spanish
  ) |>
  clean_names() |>
  mutate(gramm = ifelse(gramm == "0", "ungramm", "gramm")) |>
  mutate_if(is.character, as_factor) |> # all character variables as factors
```

```
droplevels() |>
filter(adv_type == "Deic")
```

0.1 Set contrasts

```
contrasts(df_biondo$verb_t) <- c(-0.5,+0.5)
contrasts(df_biondo$gramm) <- c(-0.5,+0.5)
contrasts(df_biondo$adv_type) <- c(-0.5,+0.5)
```

```
contrasts(df_biondo$verb_t)
```

```
      [,1]
Past    -0.5
Future   0.5
```

```
contrasts(df_biondo$gramm)
```

```
      [,1]
gramm    -0.5
ungramm   0.5
```

```
contrasts(df_biondo$adv_type)
```

```
      [,1]
Deic    -0.5
Non-deic 0.5
```

0.2 Run models

- random-intercepts only

```
fit_fp_1 <-
  lmer(log(fp) ~ verb_t*gramm +
        (1 |sj) +
        (1|item),
        data = df_biondo,
```

```
subset = roi == 4)
```

- by-item varying tense slopes

```
fit_fp_item <-
  lmerTest::lmer(log(fp) ~ verb_t*gramm +
    (1 |sj) +
    (1 + verb_t|item),
    data = df_biondo,
    subset = roi == 4)
```

1 Pooling

- do the random effects represent the exact average of participants?
 - below we see the mean logged first-pass reading time per participant (**mean**) and the by-participant intercepts from `fit_fp_1` and `fit_fp_item`
- to understand what's happening, we first have to understand pooling

```
sum_shrinkage <- df_biondo |>
  filter(roi == 4) |>
  summarise(mean = mean(log(fp), na.rm = T),
    .by = "sj") |>
  mutate(population_mean = mean(mean, na.rm = T)) |>
  left_join(coef(fit_fp_1)$sj["(Intercept)"] |> rownames_to_column(var = "sj")) |>
  rename(intercept_1 = `(Intercept)`) |>
  left_join(coef(fit_fp_item)$sj["(Intercept)"] |> rownames_to_column(var = "sj")) |>
  rename(intercept_item = `(Intercept)`)

sum_shrinkage |>
  head()
```

A tibble: 6 x 5

	sj	mean	population_mean	intercept_1	intercept_item
	<chr>	<dbl>	<dbl>	<dbl>	<dbl>
1	1	6.42	5.96	6.40	6.40
2	2	5.79	5.96	5.79	5.80
3	07	5.87	5.96	5.87	5.87
4	09	5.78	5.96	5.78	5.78
5	10	6.67	5.96	6.62	6.62

6	11	5.91	5.96	5.91	5.92
---	----	------	------	------	------

1.1 No pooling

- no pooling refers to separate regression lines fit e.g., per participant
 - each regression line is fit ignoring the population-level information
 - the intercepts are the true mean from each participant

```
head(df_no_pooling)
```

	model	sj	intercept	verb_t1	gramm1	verb_t1:gramm1
1	No pooling	1	6.422811	0.16094962	0.07844247	0.12950513
2	No pooling	2	5.792669	0.10115512	-0.10571656	-0.23199316
3	No pooling	07	5.870556	0.15344172	-0.25264603	-0.29866189
4	No pooling	09	5.780839	0.16938275	0.14074977	-0.07324559
5	No pooling	10	6.664530	0.04786447	-0.13824470	0.21824110
6	No pooling	11	5.912309	0.07573670	-0.06469794	0.35318406

```
sum_shrinkage |> head(6)
```

```
# A tibble: 6 x 5
```

	sj	mean	population_mean	intercept_1	intercept_item
	<chr>	<dbl>	<dbl>	<dbl>	<dbl>
1	1	6.42	5.96	6.40	6.40
2	2	5.79	5.96	5.79	5.80
3	07	5.87	5.96	5.87	5.87
4	09	5.78	5.96	5.78	5.78
5	10	6.67	5.96	6.62	6.62
6	11	5.91	5.96	5.91	5.92

1.2 Complete pooling

- complete pooling refers to ignoring grouping factors
 - i.e., fixed-effects only models (e.g., with `lm()` or `glm()`)
 - one regression line fit ignoring the individual-level information
 - the intercepts are the same as the population-level mean

```
head(df_pooled)
```

```
# A tibble: 6 x 6
  model      sj intercept verb_t1 gramm1 `verb_t1:gramm1`
  <chr>    <fct>    <dbl>   <dbl>   <dbl>         <dbl>
1 Complete pooling 1      5.96  0.0612 0.00310      -0.0152
2 Complete pooling 2      5.96  0.0612 0.00310      -0.0152
3 Complete pooling 07     5.96  0.0612 0.00310      -0.0152
4 Complete pooling 09     5.96  0.0612 0.00310      -0.0152
5 Complete pooling 10     5.96  0.0612 0.00310      -0.0152
6 Complete pooling 11     5.96  0.0612 0.00310      -0.0152
```

```
sum_shrinkage |> head(6)
```

```
# A tibble: 6 x 5
  sj      mean population_mean intercept_1 intercept_item
  <chr> <dbl>         <dbl>         <dbl>         <dbl>
1 1      6.42           5.96           6.40           6.40
2 2      5.79           5.96           5.79           5.80
3 07     5.87           5.96           5.87           5.87
4 09     5.78           5.96           5.78           5.78
5 10     6.67           5.96           6.62           6.62
6 11     5.91           5.96           5.91           5.92
```

1.3 Complete vs. no pooling

- complete pooling (green solid line) and no pooling (orange dotted line) of grammaticality effects for 10 participants
 - describe what you see in terms of intercept and slopes across the participants

1.4 Partial pooling: mixed models

2 Shrinkage

- turns out the estimates are pulled towards the population-level estimates
 - all the information in the model is taken into account when fitting varying intercepts and slopes

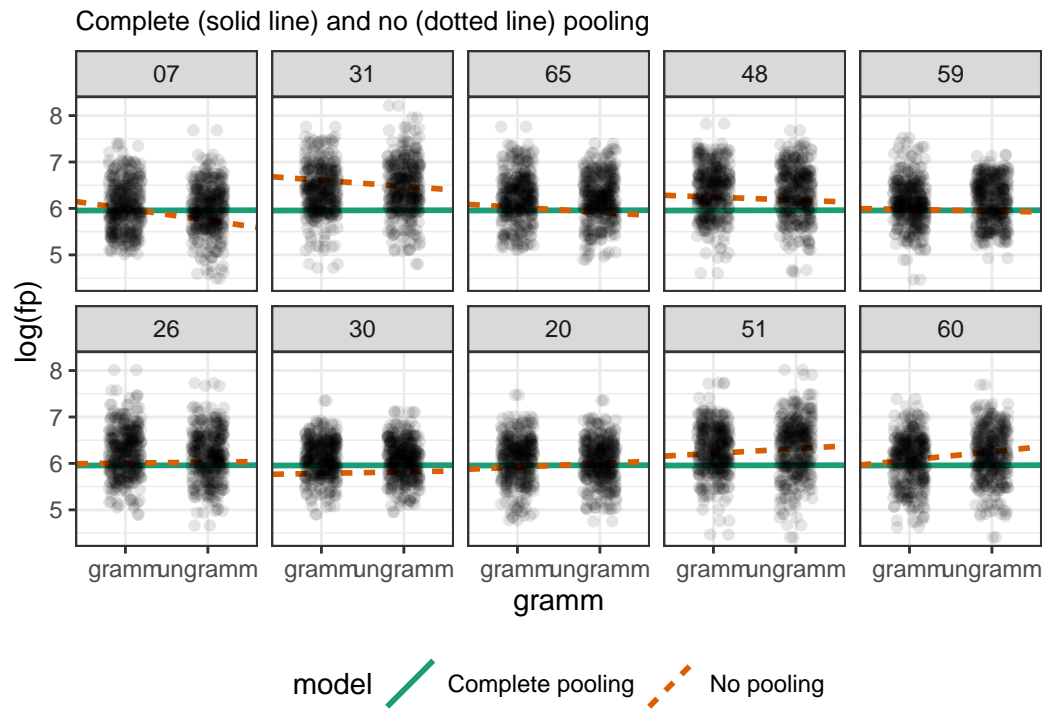


Figure 1: Observations (black dots) with complete pooling regression line (solid green) and no pooling line (dotted orange) per 10 participants

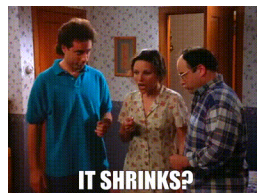


Figure 2: Elaine Benes learns about shrinkage of random effect estimates towards the population-level estimates

2.1 Shrinkage

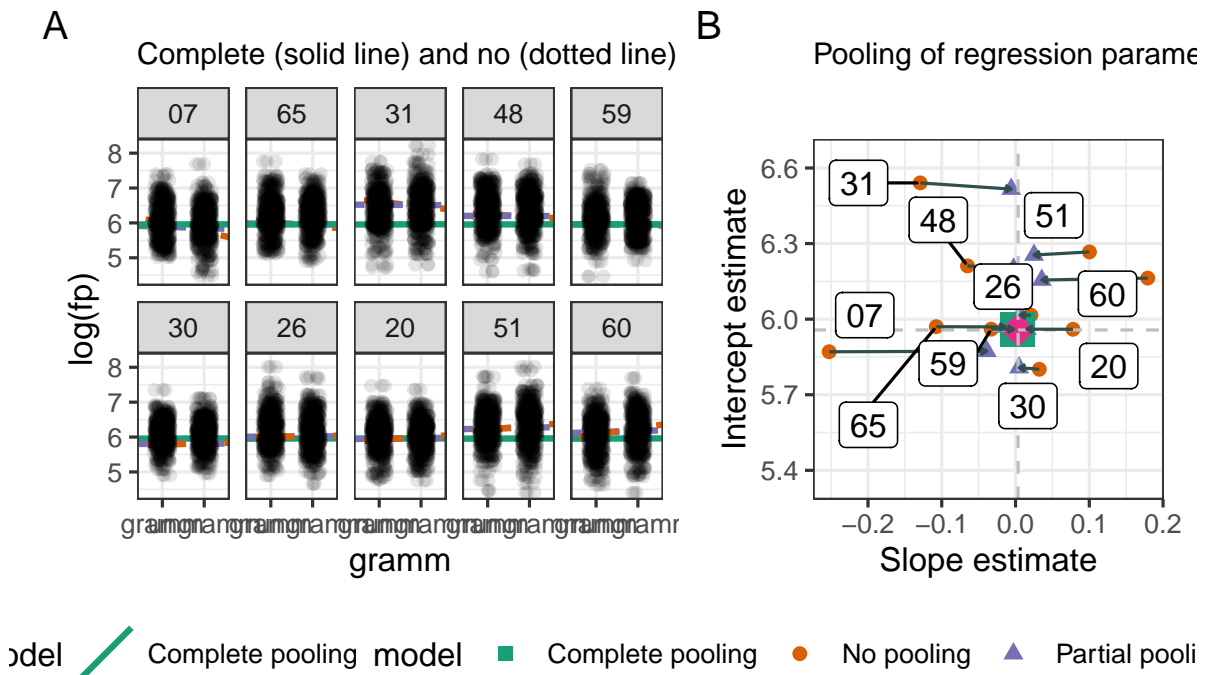


Figure 3: Shrinkage of 10 participants

2.2 Centre of gravity

- why are some points not being pulled directly to the ‘centre of gravity’?
 - they’re being pulled to a higher confidence region

3 Why shrinkage?

- with partial pooling, each random effect is like a weighted average
 - it takes into account the effect for one group level (e.g., one participant) *and* the population-level estimates
 - the empirical effect for a group level is weighted by the number of observations
 - so if one participant has fewer observations than another, then more weight is given to the population-level estimates, and vice versa
- the implications (benefits) of this:

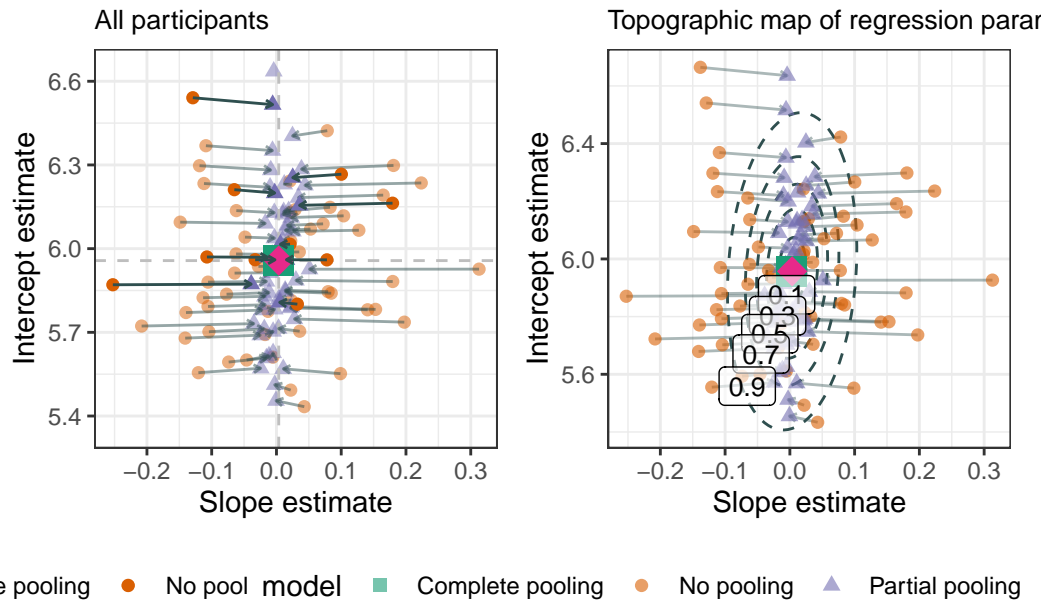


Figure 4: Shrinkage for all participants: each ellipsis represents a confidence level (really, a quantile: q1, q3, q5, q7, and q9); The inner ellipsis contains the centre 10% of the data, the outer ellipsis 90%

- imbalanced data are not a problem for linear mixed models
- the model can make predictions for unseen levels, i.e., it can generalise to new data

Learning objectives

Today we learned...

- what linear mixed models are
- how to fit a random-intercepts model
- how to inspect and interpret a mixed effects model

Important terms

Term	Definition	Equation/Code
linear mixed (effects) model	NA	NA

References

- Biondo, N., Soilemezidi, M., & Mancini, S. (2022). Yesterday is history, tomorrow is a mystery: An eye-tracking investigation of the processing of past and future time reference during sentence reading. *Journal of Experimental Psychology: Learning, Memory, and Cognition*, 48(7), 1001–1018. <https://doi.org/10.1037/xlm0001053>
- Sonderegger, M. (2023). *Regression Modeling for Linguistic Data*.
- Winter, B. (2019). Statistics for Linguists: An Introduction Using R. In *Statistics for Linguists: An Introduction Using R*. Routledge. <https://doi.org/10.4324/9781315165547>