

# RECSM: Quantitative Methods in Social Research

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# 1 Preface

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This course is about building strong quantitative social science foundations, so new AI technologies become an asset, not a black box.

This book accompanies the short course *RECSM: Quantitative Methods in Social Research*. It extends the slide decks with worked examples, code, and practice exercises that use the **European Social Survey (ESS)** microdata for the United Kingdom (GB), Germany (DE), and France (FR).

## How to use this book

- Each chapter mirrors a teaching day: descriptive statistics (Day 1), understanding associations (Day 2), nonlinear models and interactions (Day 3).
- Every exercise includes tidy, commented R code using the bundled `ess.csv` file and its HTML codebook (`ESS...subset codebook.html`).

- Required R packages are listed at the start of each chapter; install once and re-use.

If you spot anything unclear or errors, please get in touch with me.

## 2 Getting Ready

This short setup chapter ensures everyone can run the examples on their own laptop before class.

### 2.1 Install R and RStudio

- Download R from <https://cran.r-project.org/> (any recent 4.x build).
- Download RStudio Desktop from <https://posit.co/download/rstudio-desktop/> (free version).

### 2.2 Folder structure

Place the course folder anywhere convenient. The book assumes the working directory is the course root (where `ess.csv` lives). To set it inside RStudio: *Session > Set Working Directory > To Source File Location*.

```
# check current working directory
getwd()
# list course files
list.files()
```

### 2.3 Load the ESS data once

We use a pre-cleaned CSV with 80k+ respondents from GB, DE, and FR. The code below keeps only the variables used in the book and handles common missing codes (“”, 7x, 8x, 9x often mean non-response in ESS).

```
library(dplyr)
library(readr)

ess_raw <- read_csv("ess.csv", show_col_types = FALSE)

ess <- ess_raw |>
  filter(cntry %in% c("GB", "DE", "FR")) |>
  mutate(across(everything(), ~ na_if(.x, ""))) |>
  mutate(across(where(is.character), readr::parse_number, na = ""))

# quick glimpse
ess |> select(cntry, agea, gndr, ppltrst, netustm, nwsptot) |> slice_head(n = 5)
```

Tip: keep `ess` in your environment while you work across chapters to avoid reloading.

## 3 Day 1 — Describing the ESS sample

Goal: practice measurement levels, univariate summaries, and basic visualisations using the ESS subset (GB, DE, FR).

### 3.1 Variables we use

- `ppltrst` (0–10): generalised social trust (higher = more trust)
- `agea`: age in years
- `gndr`: 1 = male, 2 = female (other codes = missing)
- `nwsptot`: days per week reading newspapers (0–7; 66/77/88/99 = missing)
- `netustm`: minutes per day on the internet
- `domicil`: 1 big city ... 5 farm/countryside
- `cntry`: GB, DE, FR

```
library(dplyr)
library(ggplot2)

ess <- clean_ess()
```

### 3.2 Measurement checks

- `ppltrst` and `agea`: interval/ratio (mean and sd are fine).
- `nwsptot`, `netustm`: count-like; treat as interval for summaries but plot distributions.
- `domicil`: ordered categorical (use medians/percentiles).
- `gndr`: binary factor.

### 3.3 From description to inference

- **Descriptive vs causal inference:** describing “what is” (population levels, group gaps) vs “why” (causal claims require design/identification). Today we stay descriptive but prep for causal thinking.
- **Sampling model:** estimates come from a sample → always uncertainty. Sampling distributions tell us how much an estimator would vary across repeated samples.
- **CLT intuition:** for many statistics (like the mean), repeated samples stack up in an approximately normal bell curve as  $n$  grows; this justifies SEs, CIs, and t-tests.
- **Hypothesis testing basics:** state  $H_0/H_A$ , pick a test statistic, get a p-value, draw a conclusion while minding Type I (false positive) and Type II (false negative) risks.

### 3.4 Distributions and where centre matters

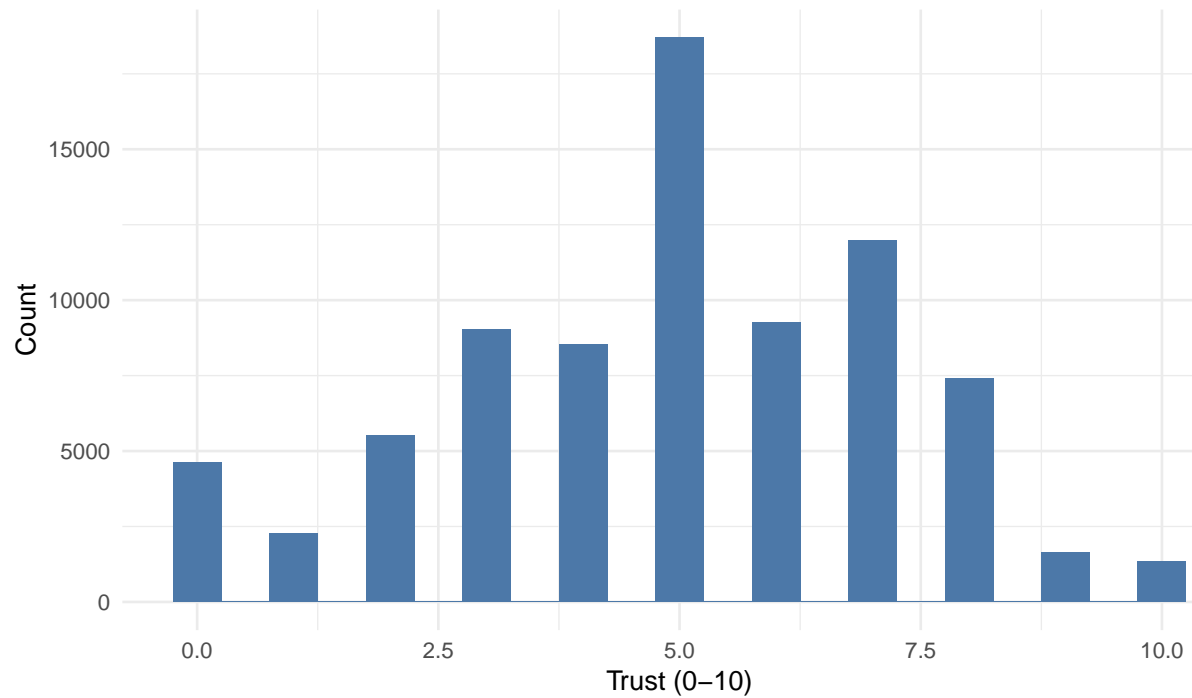
- **Central tendency:** use mean for roughly symmetric interval data (e.g., `ppltrst`), median for skewed or count-like data (e.g., `nwsptot`), and mode/proportions for categorical (`gndr`, `domicil`).
- **Shapes to look for:** symmetry vs. skew, heavy tails, spikes at 0, and multimodality. Use density/Histogram to diagnose before choosing a summary.
- **Robustness:** median and IQR resist outliers; mean and SD are more efficient when the distribution is near normal.

#### 3.4.1 Quick distribution gallery

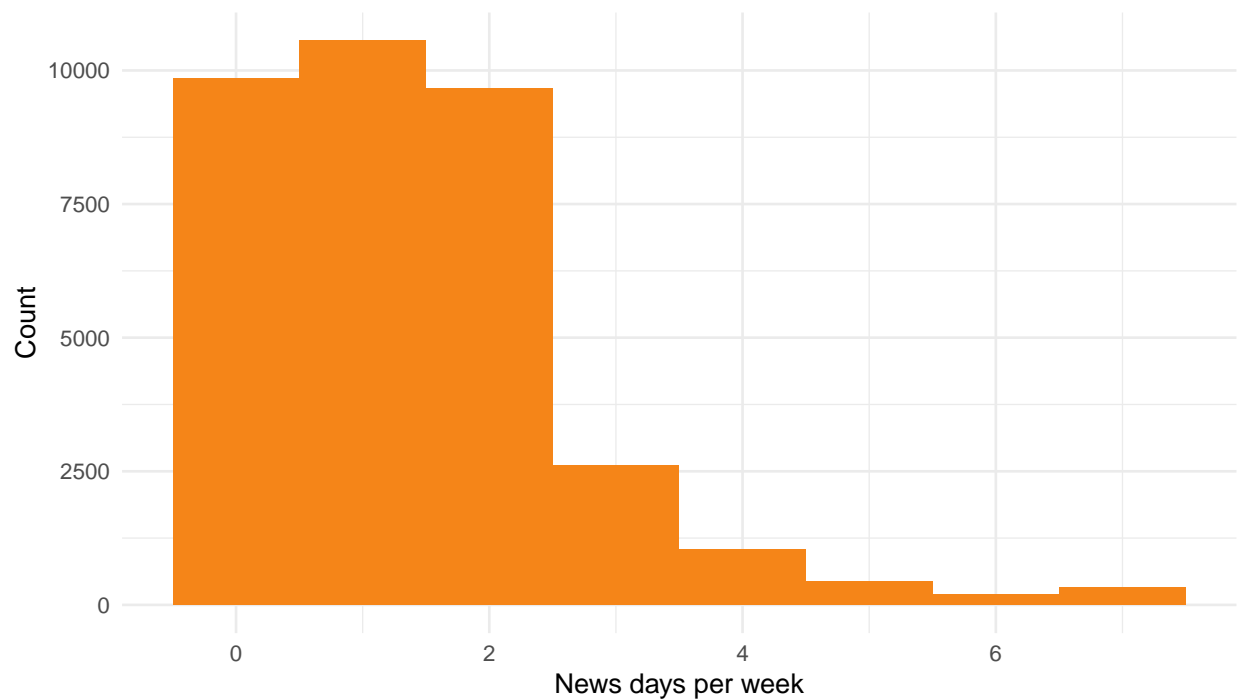
```
ess <- clean_ess()
hist_trust <- ggplot(ess, aes(x = ppltrst)) + geom_histogram(binwidth = 0.5, fill = "#4C78A8") +
  labs(x = "Trust (0-10)", y = "Count") + theme_minimal()
hist_news <- ggplot(ess, aes(x = nwsptot)) + geom_histogram(binwidth = 1, fill = "#F58518") +
```

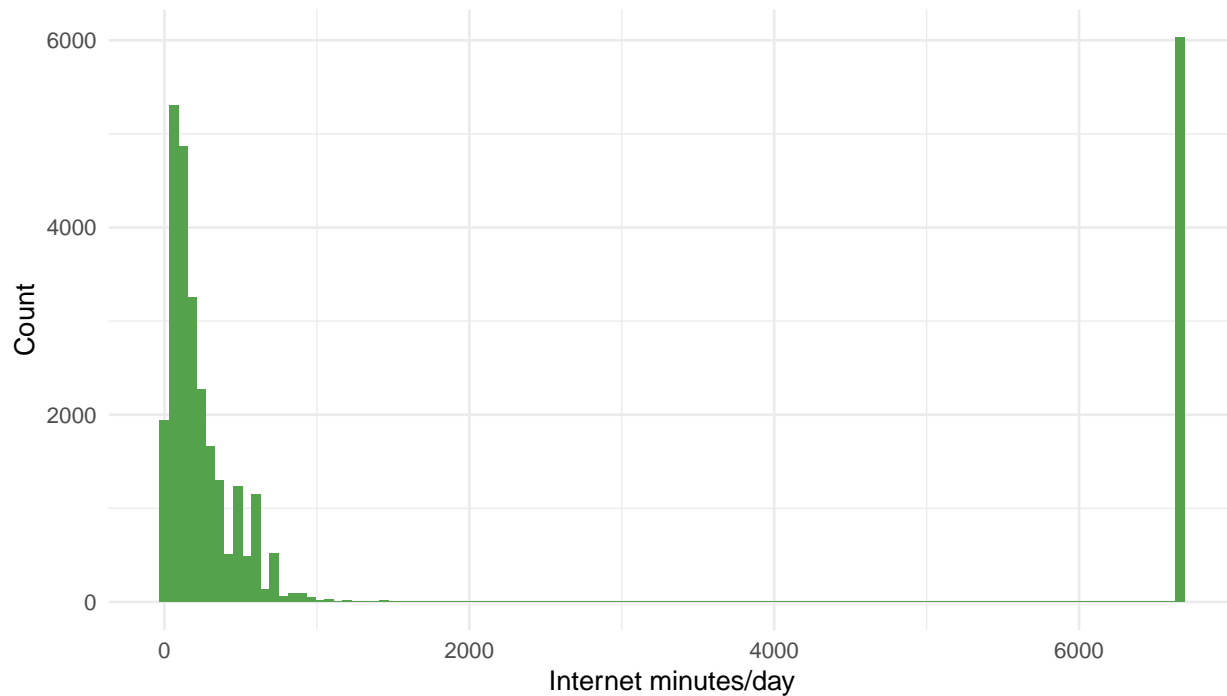
```
labs(x = "News days per week", y = "Count") + theme_minimal()
hist_net <- ggplot(ess, aes(x = netustm)) + geom_histogram(binwidth = 60, fill = "#54A24B") +
labs(x = "Internet minutes/day", y = "Count") + theme_minimal()
```

### 3.4.1.1 Code



### 3.4.1.2 Output





### 3.5 Problem set A — Univariate summaries

1. Compute mean, median, variance, and IQR for `ppltrst`, `nwsptot`, and `netustm` for each country.
2. Plot histograms (or density plots) of `ppltrst` by country; compare centres and spread.
3. Produce a table of counts and proportions for `gnr` and `domicil` by country.

#### 3.5.1 Worked example A

```
# code only (not executed in this tab)
summary_tbl <- ess |>
  group_by(cntry) |>
  summarise(
    trust_mean = mean(ppltrst, na.rm = TRUE),
    trust_sd   = sd(ppltrst, na.rm = TRUE),
    news_med   = median(nwsptot, na.rm = TRUE),
    news_iqr   = IQR(nwsptot, na.rm = TRUE),
    net_mean   = mean(netustm, na.rm = TRUE),
    net_sd     = sd(netustm, na.rm = TRUE)
  )

trust_plot <- ggplot(ess, aes(x = ppltrst, fill = cntry)) +
  geom_density(alpha = 0.35) +
  labs(x = "Social trust (0-10)", y = "Density", fill = "Country") +
  theme_minimal()

counts <- ess |>
  mutate(dom_group = factor(domicil, levels = 1:5,
```

```

labels = c("Big city", "Suburbs", "Town", "Village", "Farm")) |>
count(cntry, gender, dom_group, name = "n") |>
group_by(cntry) |>
mutate(prop = n / sum(n))

```

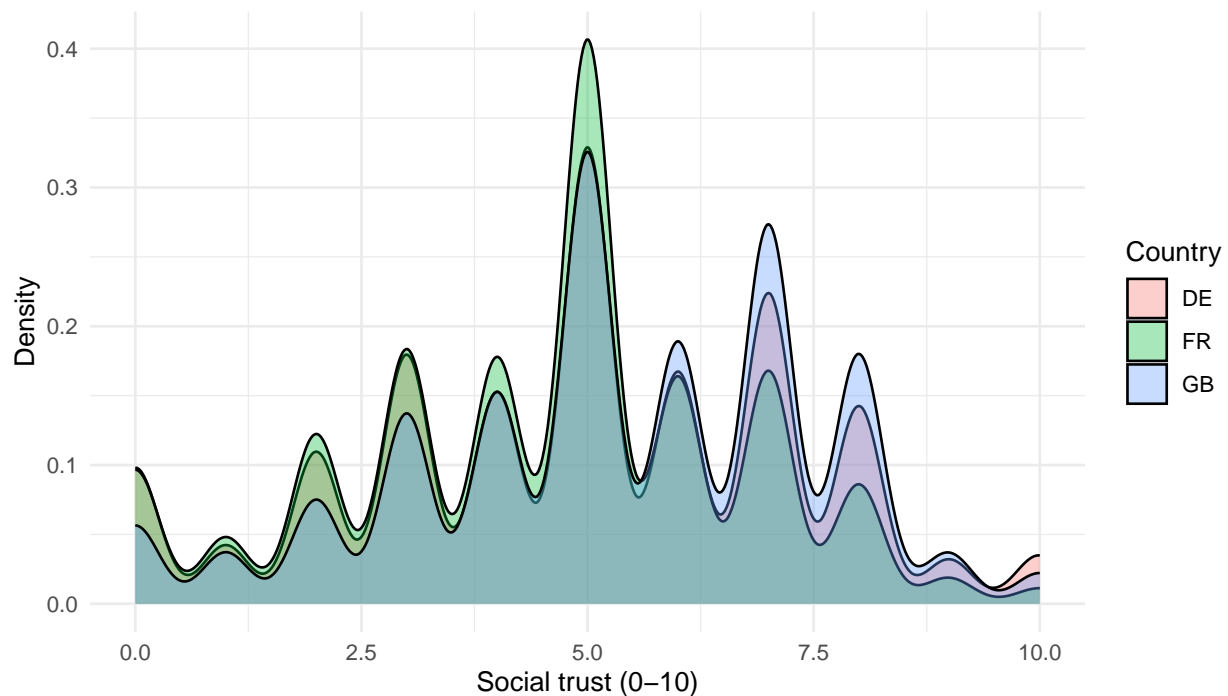
### 3.5.1.1 Code

### 3.5.1.2 Output

```

## # A tibble: 3 x 7
##   cntry trust_mean trust_sd news_med news_iqr net_mean net_sd
##   <chr>      <dbl>    <dbl>    <dbl>    <dbl>    <dbl>  <dbl>
## 1 DE         4.90      2.40        1        1    1322.  2417.
## 2 FR         4.53      2.18        1        2    1754.  2771.
## 3 GB         5.31      2.22        1        2    1539.  2581.

```



```

## # A tibble: 43 x 5
## # Groups:   cntry [3]
##   cntry gender dom_group      n  prop
##   <chr> <chr>   <fct>    <int> <dbl>
## 1 DE   Female Big city    3212 0.0872
## 2 DE   Female Suburbs   2442 0.0663
## 3 DE   Female Town      6723 0.182
## 4 DE   Female Village   5354 0.145
## 5 DE   Female Farm       391 0.0106
## 6 DE   Female <NA>        86 0.00233
## 7 DE   Male   Big city    3144 0.0853
## 8 DE   Male   Suburbs   2396 0.0650
## 9 DE   Male   Town      6475 0.176

```

```
## 10 DE      Male   Village    5739 0.156
## # i 33 more rows
```

### 3.6 Problem set B — Bivariate exploration

1. Correlate `ppltrst` with `agea` overall and by country. Does trust rise or fall with age?
2. Create side-by-side boxplots of `ppltrst` by `gender` within each country.
3. Compute the difference in mean `ppltrst` between genders; provide 95% confidence intervals using a t-test.
4. Produce a country-by-gender table of median `nwsptot`.

#### 3.6.1 Worked example B

```
# code only (not executed in this tab)
cor_age_trust <- ess |>
  group_by(cntry) |>
  summarise(corr = cor(ppltrst, agea, use = "pairwise.complete.obs"))

boxplot_trust <- ggplot(ess, aes(x = gender, y = ppltrst, fill = gender)) +
  geom_boxplot(outlier.alpha = 0.2) +
  facet_wrap(~ cntry) +
  labs(x = NULL, y = "Social trust") +
  theme_minimal() +
  theme(legend.position = "none")

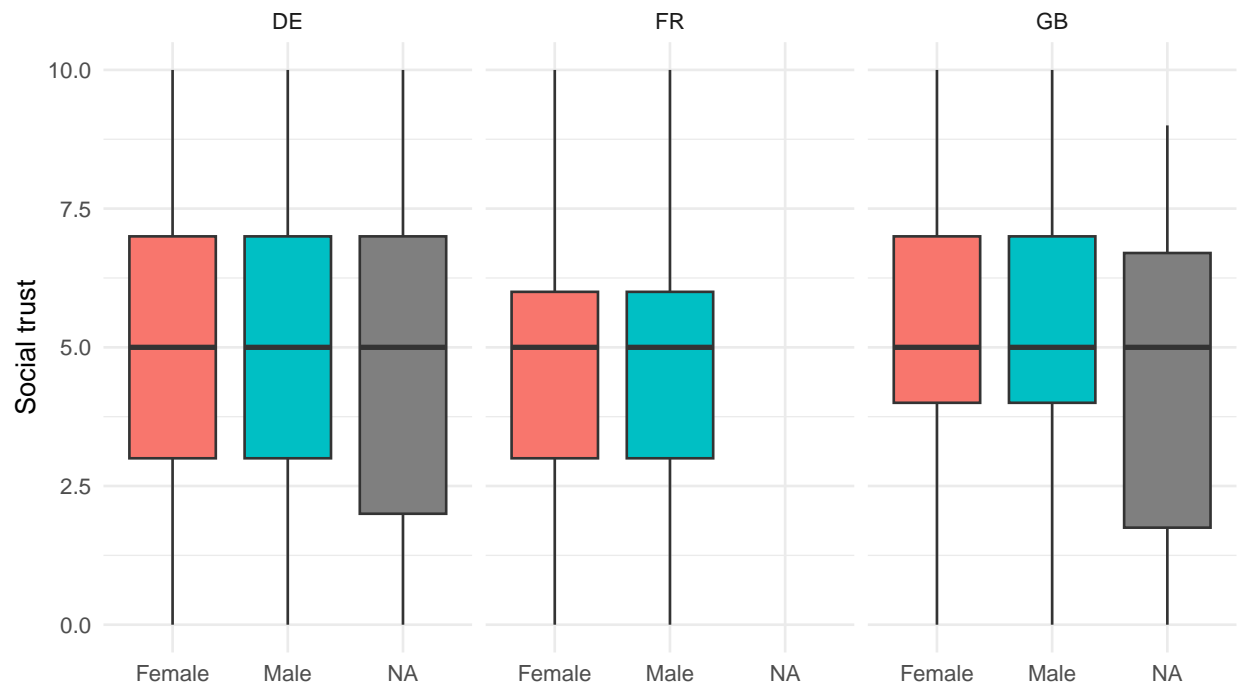
# difference in means with CI
trust_ttest <- t.test(ppltrst ~ gender, data = ess)
```

##### 3.6.1.1 Code

##### 3.6.1.2 Output

```
## # A tibble: 3 x 2
##   cntry      corr
##   <chr>    <dbl>
## 1 DE      0.00129
## 2 FR     -0.0291
## 3 GB      0.0782
```





```
##
## Welch Two Sample t-test
##
## data: ppltrst by gender
## t = -6.3769, df = 79520, p-value = 1.817e-10
## alternative hypothesis: true difference in means between group Female and group Male is not equal to
## 95 percent confidence interval:
## -0.13604975 -0.07207948
## sample estimates:
## mean in group Female    mean in group Male
##           4.870032           4.974096
```

### 3.7 Hypothesis testing quick-start

Two worked examples to introduce formal testing on Day 1.

#### 3.7.1 Example 1: Two-sample t-test (mean trust by gender)

```
t_gender <- t.test(ppltrst ~ gender, data = ess)
```

##### 3.7.1.1 Code

##### 3.7.1.2 Output

```
##
```

```
## Welch Two Sample t-test
##
## data:  ppltrst by gender
## t = -6.3769, df = 79520, p-value = 1.817e-10
## alternative hypothesis: true difference in means between group Female and group Male is not equal to
## 95 percent confidence interval:
##  -0.13604975 -0.07207948
## sample estimates:
## mean in group Female    mean in group Male
##           4.870032           4.974096
```

Interpretation: If the p-value < 0.05, we reject equal mean trust between men and women. The 95% CI shows the plausible range of the mean difference (Female – Male); if it excludes 0, the gap is statistically significant.

### 3.7.2 Example 2: Chi-squared test (gender × regular news readership)

```
ess <- ess |>
  mutate(news_regular = ifelse(nwsptot >= 3 & !nwsptot %in% c(66,77,88,99), 1, 0))

tab_news <- table(ess$gender, ess$news_regular, useNA = "no")

chi_news <- chisq.test(tab_news)
```

#### 3.7.2.1 Code

#### 3.7.2.2 Output

```
##
##           0      1
## Female 16142  2086
## Male   13917  2532

##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data:  tab_news
## X-squared = 116.47, df = 1, p-value < 2.2e-16
```

Interpretation: A small p-value means the share of regular news readers differs by gender (variables not independent). Report the chi-squared statistic, degrees of freedom, and p-value.

## 3.8 Reflection prompts

- Which variables are most skewed? How does that affect your choice of centre and spread?
- Are gender gaps in trust consistent across GB, DE, and FR?
- If `nwsptot` has many zeros, is median more informative than mean?

Use these as warm-ups before moving into regression modeling in the next chapter.

## 3.9 Sampling distributions by simulation

Central-limit intuition: as sample size grows, the sampling distribution of the mean tightens and approaches normality, even for skewed variables.

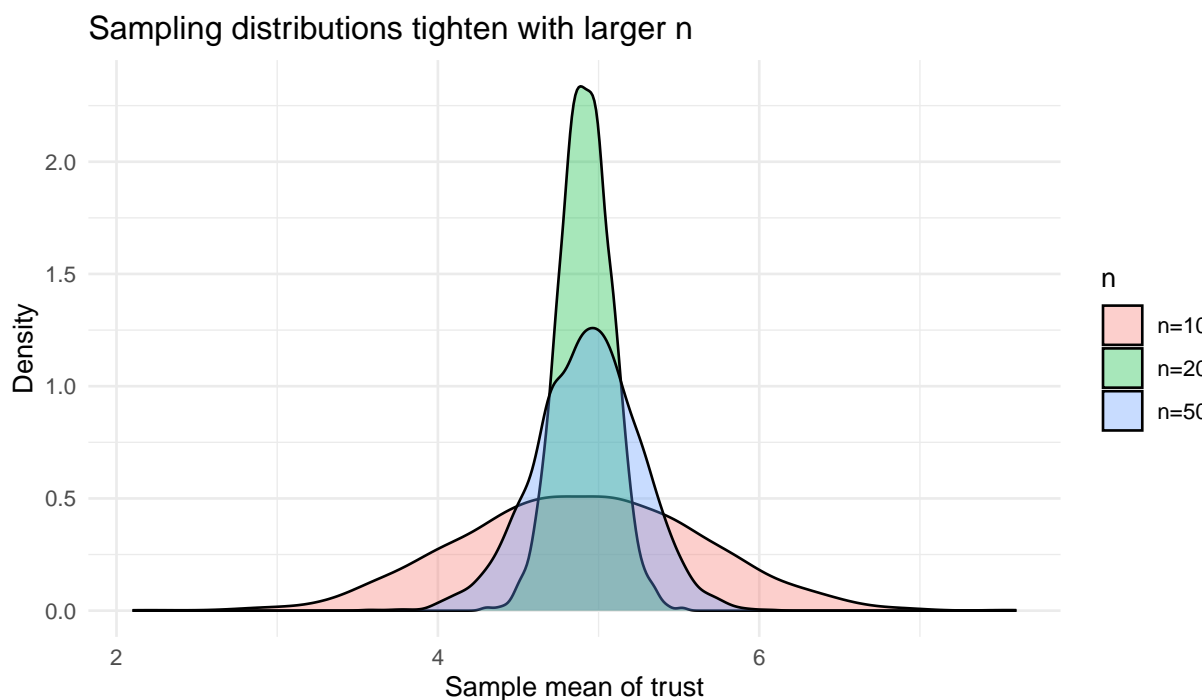
### 3.9.1 Simulating sample means of trust

```
ess <- clean_ess()
set.seed(123)

draw_means <- function(var, n, reps = 2000) {
  vals <- na.omit(var)
  replicate(reps, mean(sample(vals, n, replace = TRUE)))
}

means_n10 <- draw_means(ess$ppltrst, n = 10)
means_n50 <- draw_means(ess$ppltrst, n = 50)
means_n200 <- draw_means(ess$ppltrst, n = 200)
```

#### 3.9.1.1 Code



#### 3.9.1.2 Output

Interpretation: as  $n$  grows, the sampling distribution centers near the population mean and variance shrinks; this motivates using standard errors when reporting estimates.

### 3.10 Hypothesis testing recap (slides → practice)

- **T-tests** compare means using the t distribution (fatter tails for small  $n$ ); for two groups assume equal variances or use Welch when in doubt.
- **Chi-squared** tests independence in contingency tables.
- **Errors:**  $\alpha = P(\text{Type I})$ ,  $\beta = P(\text{Type II})$ ; lowering  $\alpha$  raises  $\beta$ . Report p-values *and* confidence intervals to show effect size and uncertainty.
- **Confidence intervals:** estimate  $\pm$  (critical value  $\times$  SE); if 95% CI excludes 0 (for mean differences), the two-sided test at  $\alpha = 0.05$  would reject  $H_0$ .

## 4 Day 2 — Linear regression with interaction effects

We replace the old dimensionality-reduction content with a deep dive on interactions. The dependent variable is **social trust** (`ppltrst`). Predictors come from media use and demographics in the ESS subset.

### Model notation recap

- Baseline linear model:  $Y_i = \beta_0 + \mathbf{x}_i^\top \beta + \varepsilon_i$ ,  $\varepsilon_i \sim \text{i.i.d. } (0, \sigma^2)$ .
- Binary–binary interaction (e.g., gender  $\times$  urban):  $Y_i = \beta_0 + \beta_1 \text{Female}_i + \beta_2 \text{Urban}_i + \beta_3 (\text{Female}_i \times \text{Urban}_i) + \dots$ 
  - $\beta_3$  is the *difference-in-differences*: the extra gap between women and men *when Urban = 1* minus the gap when Urban = 0.
- Binary–continuous interaction (gender  $\times$  age): slope for age becomes  $\beta_{\text{age}} + \beta_{\text{age} \times \text{female}} \cdot \text{Female}_i$ ; draw ribbons to see how slopes differ across groups.
- Three-way interaction (gender  $\times$  age  $\times$  country): the age slope is country- and gender-specific:  $\partial Y / \partial \text{age} = \beta_{\text{age}} + \beta_{\text{age} \times g} g + \beta_{\text{age} \times c} c + \beta_{\text{age} \times g \times c} g c$ .

```
library(dplyr)
library(ggplot2)
library(broom)
library(purrr)
library(tidyr)

source("R/clean_ess.R")

ess <- clean_ess()

# Fit once and reuse
m0 <- lm(ppltrst ~ agea + gender + news_days + country, data = ess)
m1 <- lm(ppltrst ~ gender * urban + agea + news_days + country, data = ess)
m2 <- lm(ppltrst ~ gender * agea + news_days + country, data = ess)
m3 <- lm(ppltrst ~ agea * news_days + gender + country, data = ess)
m4 <- lm(ppltrst ~ gender * agea * country + news_days, data = ess)

# Predicted values for plots (simple grids)
nd1 <- expand.grid(
  urban = c("Urban", "Non-urban"),
  gender = c("Male", "Female"),
  agea = mean(ess$agea, na.rm = TRUE),
  news_days = mean(ess$news_days, na.rm = TRUE),
  country = "GB"
```

```

)
pred1 <- predict(m1, newdata = nd1, se.fit = TRUE)
nd1$fit <- pred1$fit
nd1$lo <- pred1$fit - 1.96 * pred1$se.fit
nd1$hi <- pred1$fit + 1.96 * pred1$se.fit
int_plot1 <- ggplot(nd1, aes(x = urban, y = fit, fill = gender)) +
  geom_col(position = position_dodge(width = 0.6), width = 0.5) +
  geom_errorbar(aes(ymin = lo, ymax = hi), position = position_dodge(width = 0.6), width = 0.2) +
  labs(y = "Predicted social trust", x = "Residential area") +
  theme_minimal()

age_seq <- seq(min(ess$agea, na.rm = TRUE), max(ess$agea, na.rm = TRUE), length.out = 60)
nd2 <- expand.grid(agea = age_seq, gender = c("Male", "Female"),
  news_days = mean(ess$news_days, na.rm = TRUE),
  country = "GB")
pred2 <- predict(m2, newdata = nd2, se.fit = TRUE)
nd2$fit <- pred2$fit
nd2$lo <- pred2$fit - 1.96 * pred2$se.fit
nd2$hi <- pred2$fit + 1.96 * pred2$se.fit
int_plot2 <- ggplot(nd2, aes(x = agea, y = fit, color = gender)) +
  geom_line(size = 1) +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = gender), alpha = 0.15, color = NA) +
  labs(y = "Predicted social trust", x = "Age") +
  theme_minimal()

news_seq <- seq(min(ess$news_days, na.rm = TRUE), max(ess$news_days, na.rm = TRUE), length.out = 40)
nd3 <- expand.grid(news_days = news_seq,
  agea = quantile(ess$agea, c(.2, .5, .8), na.rm = TRUE),
  gender = "Male",
  country = "GB")
pred3 <- predict(m3, newdata = nd3, se.fit = TRUE)
nd3$fit <- pred3$fit
nd3$lo <- pred3$fit - 1.96 * pred3$se.fit
nd3$hi <- pred3$fit + 1.96 * pred3$se.fit
int_plot3 <- ggplot(nd3, aes(x = news_days, y = fit, color = factor(agea))) +
  geom_line(size = 1) +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = factor(agea)), alpha = 0.12, color = NA) +
  labs(y = "Predicted social trust", x = "News days per week", color = "Age quantile") +
  theme_minimal()

nd4 <- expand.grid(agea = age_seq,
  gender = c("Male", "Female"),
  country = c("GB", "DE", "FR"),
  news_days = mean(ess$news_days, na.rm = TRUE))
pred4 <- predict(m4, newdata = nd4, se.fit = TRUE)
nd4$fit <- pred4$fit
nd4$lo <- pred4$fit - 1.96 * pred4$se.fit
nd4$hi <- pred4$fit + 1.96 * pred4$se.fit
int_plot4 <- ggplot(nd4, aes(x = agea, y = fit, color = gender)) +
  geom_line() +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = gender), alpha = 0.12, color = NA) +
  facet_wrap(~ country) +
  labs(y = "Predicted social trust", x = "Age") +

```

```

theme_minimal()

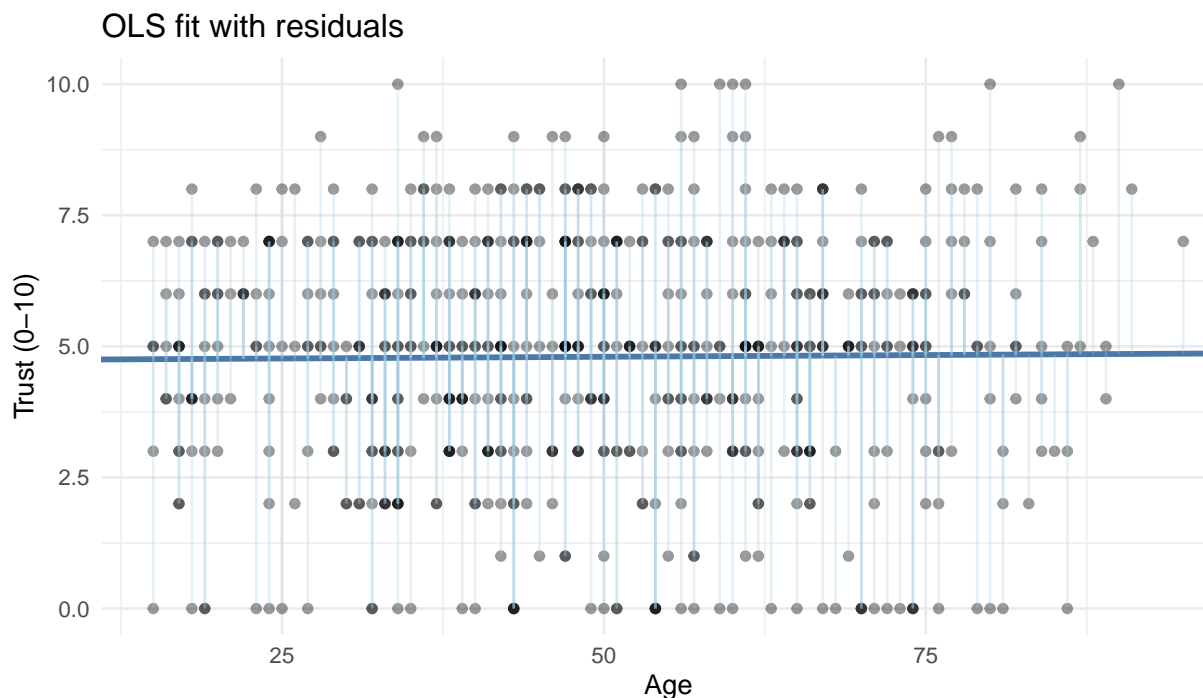
# OLS intuition demo (interactive)
ess_small <- ess |> select(ppltrst, agea) |> drop_na() |> slice_sample(n = 600)
fit_simple <- lm(ppltrst ~ agea, data = ess_small)

base_resid <- ggplot(ess_small, aes(x = agea, y = pptrst)) +
  geom_point(alpha = 0.4) +
  geom_abline(slope = coef(fit_simple)[2], intercept = coef(fit_simple)[1], color = "#4C78A8", size = 1) +
  geom_segment(aes(xend = agea, yend = fitted(fit_simple)), alpha = 0.25, color = "#9ecae1") +
  labs(x = "Age", y = "Trust (0-10)", title = "OLS fit with residuals") +
  theme_minimal()

slope_grid <- seq(coef(fit_simple)[2] - 0.08, coef(fit_simple)[2] + 0.08, length.out = 30)
anim_df <- map_dfr(slope_grid, ~{
  pred <- coef(fit_simple)[1] + .x * ess_small$agea
  tibble(agea = ess_small$agea,
    pptrst = ess_small$ppltrst,
    slope = sprintf("%.3f", .x),
    pred = pred,
    resid = pptrst - pred)
})
anim_plot <- ggplot(anim_df, aes(x = agea, y = pptrst, frame = slope)) +
  geom_point(alpha = 0.35) +
  geom_abline(aes(slope = as.numeric(slope), intercept = coef(fit_simple)[1]), color = "#4C78A8", size = 1) +
  labs(x = "Age", y = "Trust (0-10)", title = "Searching for the best-fit slope") +
  theme_minimal()

```

## 4.1 OLS intuition: best-fit line and residuals



### 4.1.0.1 Output

Interpretation: the static panel shows residuals; if you run the optional plotly code locally, the moving line illustrates how residuals shrink as the slope approaches the OLS solution.

## 4.2 1. Baseline linear model

```
m0 <- lm(ppltrst ~ agea + gender + news_days + country, data = ess)
broom::tidy(m0)
```

### 4.2.0.1 Code

### 4.2.0.2 Output

```
## # A tibble: 6 x 5
##   term          estimate std.error statistic  p.value
##   <chr>          <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)   4.68      0.0394    119.      0
## 2 agea         -0.00245   0.000691   -3.54 4.01e- 4
## 3 genderMale    0.101     0.0246     4.12 3.84e- 5
## 4 news_days     0.0727    0.00988     7.36 1.93e-13
## 5 countryFR    -0.244     0.0307    -7.94 2.00e-15
## 6 countryGB     0.545     0.0287    19.0 9.60e-80
```

Interpretation: Trust increases slightly with age and differs by country and gender; focus on sign and magnitude of the coefficients rather than raw p-values when discussing effect sizes.

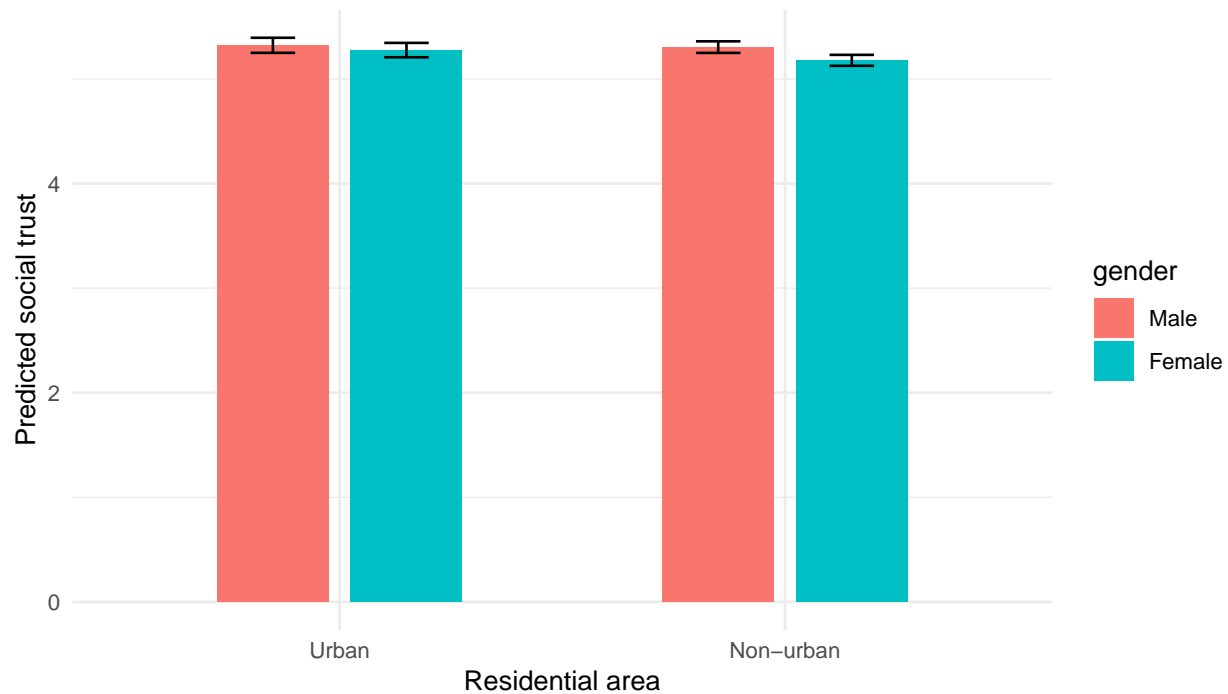
## 4.3 2. Binary $\times$ Binary interaction (gender $\times$ urban)

```
m1 <- lm(ppltrst ~ gender * urban + agea + news_days + country, data = ess)
int_plot1 <- ggplot(nd1, aes(x = urban, y = fit, fill = gender)) +
  geom_col(position = position_dodge(width = 0.6), width = 0.5) +
  labs(y = "Predicted social trust", x = "Residential area") +
  theme_minimal()
```

### 4.3.0.1 Code

### 4.3.0.2 Output

```
## # A tibble: 8 x 5
##   term                estimate std.error statistic  p.value
##   <chr>                <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)         4.65      0.0412    113.      0
## 2 genderMale          0.126     0.0295     4.28 1.85e- 5
## 3 urbanUrban          0.0979    0.0365     2.68 7.34e- 3
## 4 agea               -0.00238   0.000692   -3.44 5.92e- 4
## 5 news_days           0.0716    0.00990     7.23 4.79e-13
## 6 countryFR          -0.244     0.0307    -7.96 1.71e-15
## 7 countryGB           0.546     0.0288    19.0 5.96e-80
## 8 genderMale:urbanUrban -0.0806   0.0529    -1.52 1.28e- 1
```



Interpretation: The urban–rural trust gap is small; note whether the CI bars for Male vs Female overlap. If they do, the moderation by gender is likely negligible.

Interpretation focus: Does the urban–rural gap differ by gender?

#### 4.4 3. Binary $\times$ Continuous interaction (gender $\times$ age)

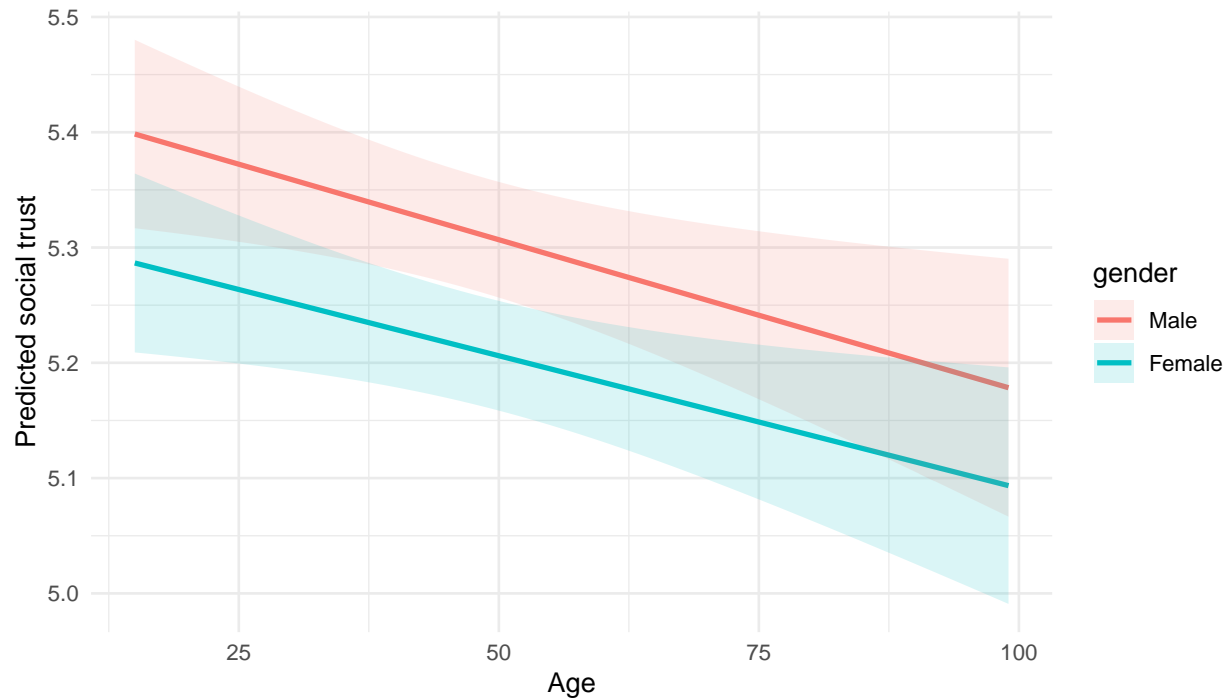
```
m2 <- lm(ppltrst ~ gender * agea + news_days + country, data = ess)
int_plot2 <- ggplot(nd2, aes(x = agea, y = fit, color = gender)) +
  geom_line(size = 1) +
  labs(y = "Predicted social trust", x = "Age") +
  theme_minimal()
```

##### 4.4.0.1 Code

##### 4.4.0.2 Output

```
## # A tibble: 7 x 5
##   term                estimate std.error statistic  p.value
##   <chr>              <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)        4.68      0.0498     93.9      0
## 2 genderMale         0.117     0.0691      1.69 9.14e- 2
## 3 agea              -0.00230  0.000925    -2.49 1.29e- 2
## 4 news_days          0.0727     0.00988     7.36 1.88e-13
## 5 countryFR         -0.244     0.0307    -7.94 2.05e-15
## 6 countryGB          0.545     0.0287    19.0 9.42e-80
## 7 genderMale:agea    -0.000321  0.00134    -0.240 8.10e- 1
```





Interpretation: Slopes by age differ by gender; parallel ribbons would imply no interaction. Diverging ribbons indicate the age effect depends on gender.

Key idea: slopes for age are estimated separately for men and women.

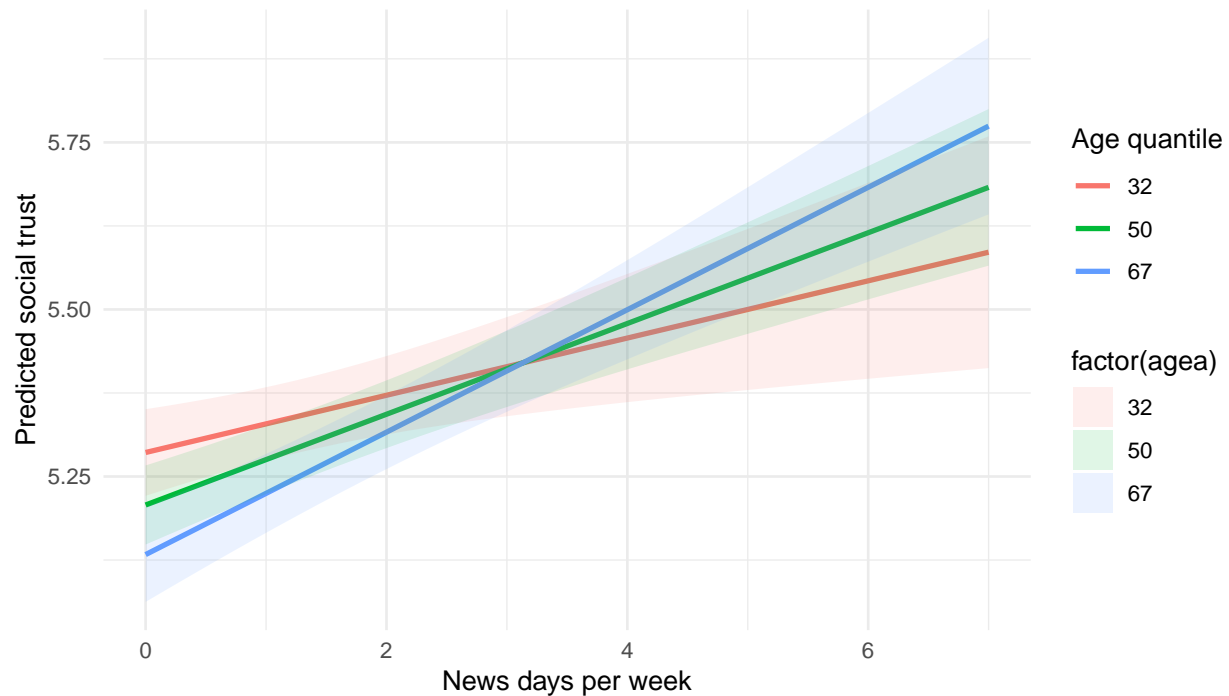
#### 4.5 4. Continuous $\times$ Continuous interaction (age $\times$ news consumption)

```
m3 <- lm(ppltrst ~ agea * news_days + gender + country, data = ess)
int_plot3 <- ggplot(nd3, aes(x = news_days, y = fit, color = factor(agea))) +
  geom_line(size = 1) +
  labs(y = "Predicted social trust", x = "News days per week", color = "Age quantile") +
  theme_minimal()
```

##### 4.5.0.1 Code

##### 4.5.0.2 Output

```
## # A tibble: 7 x 5
##   term                estimate std.error statistic  p.value
##   <chr>              <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)        4.78      0.0516     92.6      0
## 2 agea              -0.00436  0.000975    -4.48 7.65e- 6
## 3 news_days         -0.00178  0.0285     -0.0627 9.50e- 1
## 4 genderMale         0.101    0.0246      4.12 3.75e- 5
## 5 countryFR         -0.243    0.0307     -7.92 2.46e-15
## 6 countryGB         0.548    0.0288     19.1 1.57e-80
## 7 agea:news_days     0.00139  0.000500      2.79 5.31e- 3
```



Interpretation: Check whether the news-consumption slope changes across age quantiles; overlapping ribbons mean little moderation, separated ribbons suggest stronger news effects at certain ages.

Discuss whether news exposure moderates the age-trust relationship.

## 4.6 5. Three-way interaction (gender $\times$ age $\times$ country)

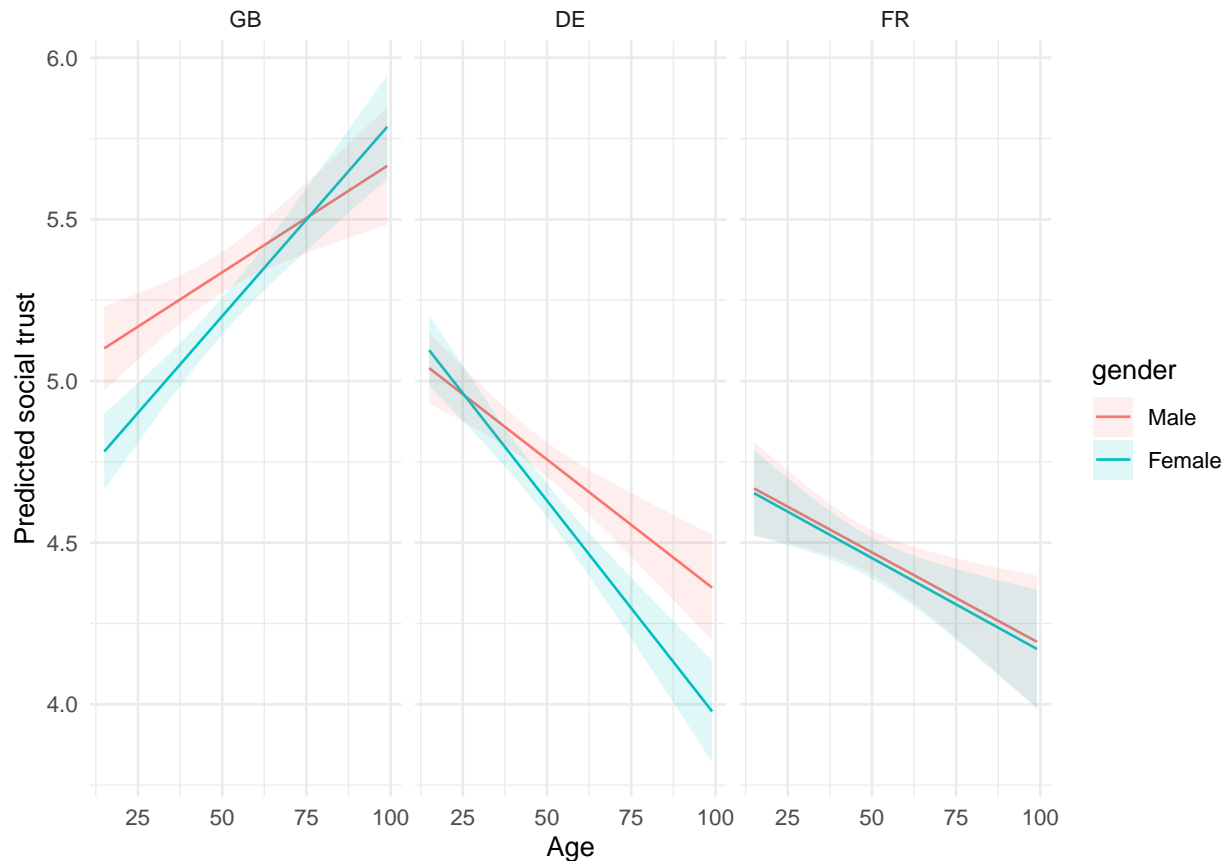
```
m4 <- lm(ppltrst ~ gender * agea * country + news_days, data = ess)
int_plot4 <- ggplot(nd4, aes(x = agea, y = fit, color = gender)) +
  geom_line() +
  facet_wrap(~ country) +
  labs(y = "Predicted social trust", x = "Age") +
  theme_minimal()
```

### 4.6.0.1 Code

### 4.6.0.2 Output

```
## # A tibble: 13 x 5
##   term                                estimate std.error statistic  p.value
##   <chr>                                <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)                        5.19      0.0760     68.2      0
## 2 genderMale                       -0.133     0.107     -1.24  2.14e- 1
## 3 agea                             -0.0133    0.00149   -8.92  4.69e-19
## 4 countryFR                        -0.555     0.119     -4.69  2.79e- 6
## 5 countryGB                       -0.692     0.111     -6.25  4.24e-10
## 6 news_days                         0.0775    0.00988    7.84  4.66e-15
## 7 genderMale:agea                   0.00521    0.00211    2.48  1.33e- 2
```

```
## 8 genderMale:countryFR      0.147      0.173      0.849 3.96e- 1
## 9 genderMale:countryGB      0.531      0.162      3.28  1.04e- 3
## 10 agea:countryFR           0.00756    0.00229      3.30  9.58e- 4
## 11 agea:countryGB           0.0253     0.00213     11.9  1.92e-32
## 12 genderMale:agea:countryFR -0.00512   0.00335     -1.53  1.26e- 1
## 13 genderMale:agea:countryGB -0.0104    0.00312     -3.35  8.23e- 4
```



Interpretation: Three-way plots show country-specific age slopes by gender; look for countries where ribbons separate widely—that’s where the interaction is substantive.

Strategy: interpret pairwise contrasts within each country before comparing across countries.

## 4.7 6. Model comparison and diagnostics

```
broom::glance(m0, m1, m2, m3, m4)
# quick residual check
par(mfrow = c(2,2)); plot(m3)
```

## 4.8 Problem set — Interaction lab

1. Refit `m1` but swap `urban` with a binary indicator for *high education* (e.g., `edu yrs >= 15`). Interpret the gender gap at low vs high education.
2. Build a model with `ppltrst ~ news_days * country + agea + gender`. Compute marginal effects of `news_days` within each country.

3. Add a three-way term `gender * urban * country`. Plot predicted trust for all six gender-by-urban-by-country profiles.
4. Briefly report which interaction improves fit (compare adjusted  $R^2$  and AIC) and whether the effect is substantively meaningful.

Use `marginalEffects::plot_slopes()` and `plot_predictions()` to visualise interactions instead of only staring at coefficients.

## 5 Day 3 — Non-linear models: logistic regression

We now model binary outcomes. This replaces the old factor-analysis/PCA content and links three specifications:

- **Linear probability model (LPM):**  $Y_i \in \{0, 1\}$ ,  $\mathbb{E}[Y_i | X_i] = X_i\beta$  (identity link).
- **Logit:**  $p_i = \Pr(Y_i = 1 | X_i) = \text{logit}^{-1}(X_i\beta) = \frac{e^{X_i\beta}}{1+e^{X_i\beta}}$ .
- **Marginal effects:**  $\frac{\partial p_i}{\partial x_{ik}} = p_i(1-p_i)\beta_k$ , highlighting how effects vary with the baseline probability.
- **Interaction in logit:** For  $p_i = \text{logit}^{-1}(\eta_i)$  with  $\eta_i = \beta_0 + \beta_1x + \beta_2z + \beta_3xz$ , the cross-partial effect is  $\frac{\partial^2 p_i}{\partial x \partial z} = p_i(1-p_i)(1-2p_i)\beta_1\beta_2 + p_i(1-p_i)\beta_3$ ; sign can vary with  $p_i$ .

**Outcome:** `news_regular` = 1 if a respondent reads newspapers on at least 3 days per week (`nwsptot >= 3`), 0 otherwise.

**Predictors:** age (`agea`), gender (`gnr`), country (`cntry`), education (`eduyrs`), and their interactions.

```
library(dplyr)
library(broom)
library(ggplot2)

source("R/clean_ess.R")

ess <- clean_ess()

# Fit once and reuse
lpm1 <- lm(news_regular ~ agea + gender + country, data = ess)
logit1 <- glm(news_regular ~ agea + gender + country, data = ess, family = binomial())
logit2 <- glm(news_regular ~ gender * country + agea, data = ess, family = binomial())
logit3 <- glm(news_regular ~ gender * agea + country + eduyrs, data = ess, family = binomial())
logit4 <- glm(news_regular ~ agea * eduyrs + gender + country, data = ess, family = binomial())
logit5 <- glm(news_regular ~ gender * agea * country + eduyrs, data = ess, family = binomial())

# Prediction data for plots
age_seq <- seq(min(ess$agea, na.rm = TRUE), max(ess$agea, na.rm = TRUE), length.out = 60)
edu_seq <- seq(min(ess$eduyrs, na.rm = TRUE), max(ess$eduyrs, na.rm = TRUE), length.out = 40)

nd2 <- expand.grid(country = c("GB", "DE", "FR"),
                  gender = c("Male", "Female"),
                  agea = mean(ess$agea, na.rm = TRUE))
pred2 <- predict(logit2, newdata = nd2, type = "link", se.fit = TRUE)
nd2$pr <- plogis(pred2$fit)
nd2$lo <- plogis(pred2$fit - 1.96 * pred2$se.fit)
nd2$hi <- plogis(pred2$fit + 1.96 * pred2$se.fit)
```

```

plot_gender_country <- ggplot(nd2, aes(x = country, y = pr, fill = gender)) +
  geom_col(position = position_dodge(width = 0.6), width = 0.5) +
  geom_errorbar(aes(ymin = lo, ymax = hi), position = position_dodge(width = 0.6), width = 0.2) +
  labs(y = "Pr(regular news)") +
  theme_minimal()

nd3 <- expand.grid(agea = age_seq, gender = c("Male", "Female"),
                  country = "GB", eduyrs = mean(ess$eduyrs, na.rm = TRUE))
pred3 <- predict(logit3, newdata = nd3, type = "link", se.fit = TRUE)
nd3$pr <- plogis(pred3$fit)
nd3$lo <- plogis(pred3$fit - 1.96 * pred3$se.fit)
nd3$hi <- plogis(pred3$fit + 1.96 * pred3$se.fit)
plot_gender_age <- ggplot(nd3, aes(x = agea, y = pr, color = gender)) +
  geom_line(size = 1) +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = gender), alpha = 0.15, color = NA) +
  labs(y = "Pr(regular news)", x = "Age") +
  theme_minimal()

nd4 <- expand.grid(eduyrs = edu_seq,
                  agea = quantile(ess$agea, c(.2, .5, .8), na.rm = TRUE),
                  gender = "Male", country = "GB")
pred4 <- predict(logit4, newdata = nd4, type = "link", se.fit = TRUE)
nd4$pr <- plogis(pred4$fit)
nd4$lo <- plogis(pred4$fit - 1.96 * pred4$se.fit)
nd4$hi <- plogis(pred4$fit + 1.96 * pred4$se.fit)
plot_age_edu <- ggplot(nd4, aes(x = eduyrs, y = pr, color = factor(agea))) +
  geom_line(size = 1) +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = factor(agea)), alpha = 0.12, color = NA) +
  labs(y = "Pr(regular news)", x = "Years of education", color = "Age quantile") +
  theme_minimal()

nd5 <- expand.grid(agea = age_seq,
                  gender = c("Male", "Female"),
                  country = c("GB", "DE", "FR"),
                  eduyrs = mean(ess$eduyrs, na.rm = TRUE))
pred5 <- predict(logit5, newdata = nd5, type = "link", se.fit = TRUE)
nd5$pr <- plogis(pred5$fit)
nd5$lo <- plogis(pred5$fit - 1.96 * pred5$se.fit)
nd5$hi <- plogis(pred5$fit + 1.96 * pred5$se.fit)
plot_threeway <- ggplot(nd5, aes(x = agea, y = pr, color = gender)) +
  geom_line() +
  geom_ribbon(aes(ymin = lo, ymax = hi, fill = gender), alpha = 0.12, color = NA) +
  facet_wrap(~ country) +
  labs(y = "Pr(regular news)", x = "Age") +
  theme_minimal()

# Average marginal effects (finite-difference, no parallelism)
calc_ame <- function(model, data, var, step = 1) {
  data_hi <- data
  if (is.numeric(data_hi[[var]])) {
    data_hi[[var]] <- data_hi[[var]] + step
  } else if (is.factor(data_hi[[var]]) || is.character(data_hi[[var]])) {
    # flip binary factor

```

```

    if (all(na.omit(unique(data_hi[[var]])) %in% c("Male", "Female"))) {
      data_hi[[var]] <- ifelse(data_hi[[var]] == "Female", "Male", "Female")
    }
  }
  p_hi <- predict(model, newdata = data_hi, type = "response")
  p_lo <- predict(model, newdata = data, type = "response")
  mean(p_hi - p_lo, na.rm = TRUE)
}

ame_age <- calc_ame(logit5, ess, "agea", step = 1)
ame_fem <- calc_ame(logit5, ess, "gender", step = 0) # Female vs Male switch
ame_table <- tibble(
  variable = c("agea (+1 year)", "gender (Female vs Male)"),
  AME = c(ame_age, ame_fem)
)

# Marginal effect of being Female at age 25 vs 65
nd_female <- data.frame(agea = c(25, 65),
  gender = "Female",
  country = "GB",
  eduys = mean(ess$eduys, na.rm = TRUE))
nd_male <- nd_female; nd_male$gender <- "Male"
pred_f <- predict(logit5, newdata = nd_female, type = "response")
pred_m <- predict(logit5, newdata = nd_male, type = "response")
me_female_diff <- data.frame(agea = c(25, 65),
  diff = pred_f - pred_m)
me_female_plot <- ggplot(me_female_diff, aes(x = factor(agea), y = diff)) +
  geom_col(fill = "#4C78A8", width = 0.5) +
  geom_hline(yintercept = 0, linetype = "dashed") +
  labs(x = "Age", y = "Pr(Female) - Pr(Male)", title = "Marginal effect of being Female") +
  theme_minimal()

```

## 5.1 0. Linear probability model (LPM) first

- **Specification:**  $Y_i = X_i\beta + \varepsilon_i$ ,  $Y_i \in \{0, 1\}$ . Ordinary least squares with a binary outcome.
- **Pros:** Coefficients are immediate probability changes ( $\Delta p$ ) per unit of  $X$ ; easy to interpret and to add fixed effects.
- **Cons:** Predicted values can leave  $[0, 1]$ ; errors are heteroskedastic; marginal effects are assumed constant even when baseline risk is near 0 or 1.
- **When is LPM “safe enough”?** Middle-range probabilities (e.g., 0.2–0.8), modest leverage points, and when the goal is fast descriptive decomposition or fixed-effects absorption. Use robust standard errors.

```

library(sandwich)
library(lmtest)

lpm_vcov <- sandwich::vcovHC(lpm1, type = "HC1")
lpm_tidy <- broom::tidy(lpm1, conf.int = TRUE, vcov = lpm_vcov)

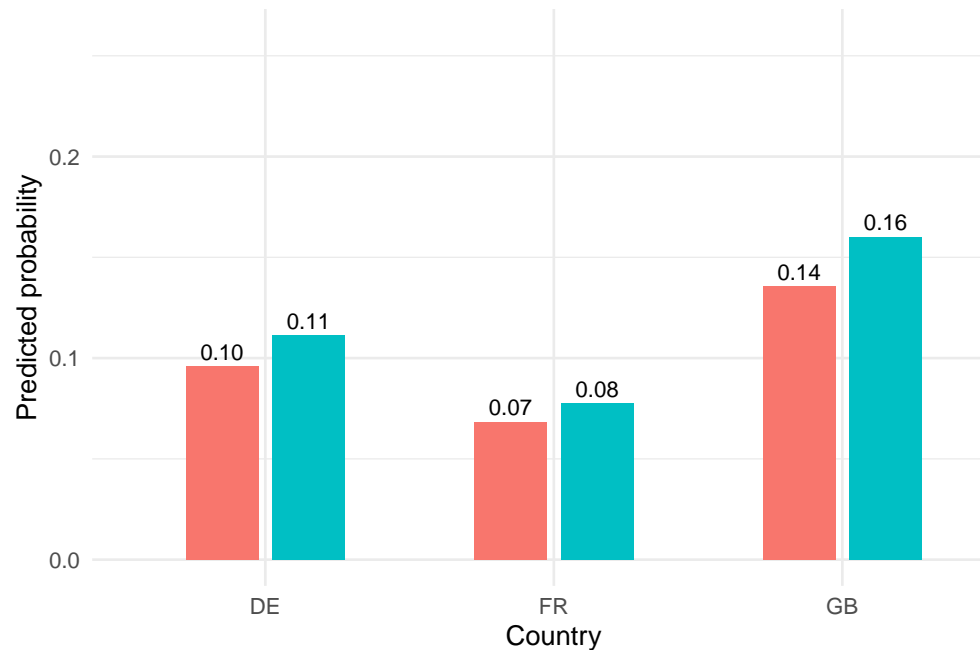
```

### 5.1.0.1 Output

```
## # A tibble: 5 x 7
```

##	term	estimate	std.error	statistic	p.value	conf.low	conf.high
##	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
## 1	(Intercept)	-0.0861	0.00569	-15.1	1.43e-51	-0.0973	-0.0750
## 2	agea	0.00397	0.0000969	40.9	0	0.00378	0.00416
## 3	genderMale	0.0429	0.00356	12.0	2.31e-33	0.0359	0.0498
## 4	countryFR	-0.0334	0.00442	-7.56	4.08e-14	-0.0421	-0.0248
## 5	countryGB	0.0490	0.00418	11.7	9.41e-32	0.0408	0.0572

LPM vs logit baseline predictions



### 5.1.0.2 LPM vs logit predictions

Observation: If LPM bars stray above 1 or below 0, that signals the need for a logit/probit link. Here the mid-range outcome keeps LPM close, but we switch to logit next for coherent probabilities and curved marginal effects.

## 5.2 1. Simple logistic model

### 5.2.1 Simple logistic model

```
logit1 <- glm(news_regular ~ agea + gender + country, data = ess, family = binomial())
broom::tidy(logit1, exponentiate = TRUE, conf.int = TRUE)
```

5.2.1.1 Code Odds ratios > 1 indicate higher odds of being a regular news reader.

### 5.2.1.2 Output

```
## # A tibble: 5 x 7
##   term      estimate std.error statistic  p.value conf.low conf.high
##   <chr>      <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
```

## 1 (Intercept)	0.0175	0.0624	-64.9	0	0.0155	0.0197
## 2 agea	1.04	0.000941	38.5	0	1.04	1.04
## 3 genderMale	1.50	0.0330	12.2	3.33e-34	1.40	1.60
## 4 countryFR	0.692	0.0452	-8.16	3.34e-16	0.633	0.755
## 5 countryGB	1.48	0.0367	10.7	9.12e-27	1.38	1.59

Interpretation: Odds ratios >1 raise the chance of regular news use; check if the CI excludes 1 for age, gender, or specific countries before claiming significance.

### 5.3 2. Binary $\times$ Binary interaction (gender $\times$ country)

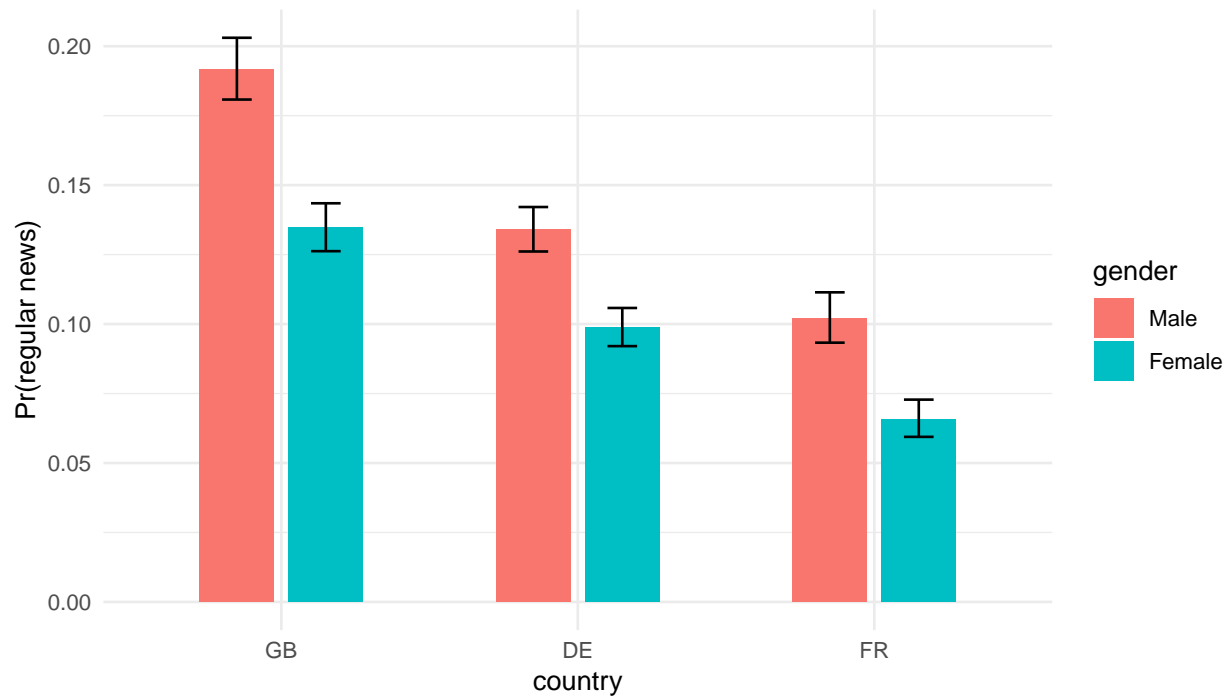
```
logit2 <- glm(news_regular ~ gender * country + agea, data = ess, family = binomial())
plot_gender_country <- ggplot(nd2, aes(x = country, y = pr, fill = gender)) +
  geom_col(position = position_dodge(width = 0.6), width = 0.5) +
  labs(y = "Pr(regular news)") +
  theme_minimal()
```

#### 5.3.0.1 Code

#### 5.3.0.2 Output

```
## # A tibble: 7 x 7
##   term                estimate std.error statistic  p.value conf.low conf.high
##   <chr>                <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)          0.0180  0.0662   -60.7    0        0.0158  0.0205
## 2 genderMale           1.41    0.0522    6.61 3.83e-11  1.27    1.56
## 3 countryFR            0.643   0.0673   -6.57 5.14e-11  0.563   0.733
## 4 countryGB            1.42    0.0536    6.55 5.87e-11  1.28    1.58
## 5 agea                 1.04    0.000941 38.5    0        1.04    1.04
## 6 genderMale:countryFR 1.14    0.0907    1.47 1.42e- 1  0.957   1.37
## 7 genderMale:countryGB 1.08    0.0735    1.04 2.97e- 1  0.935   1.25
```





Interpretation: If male–female bars overlap within countries (CI bars), gender differences are modest. Compare across countries to see where the gap is largest.

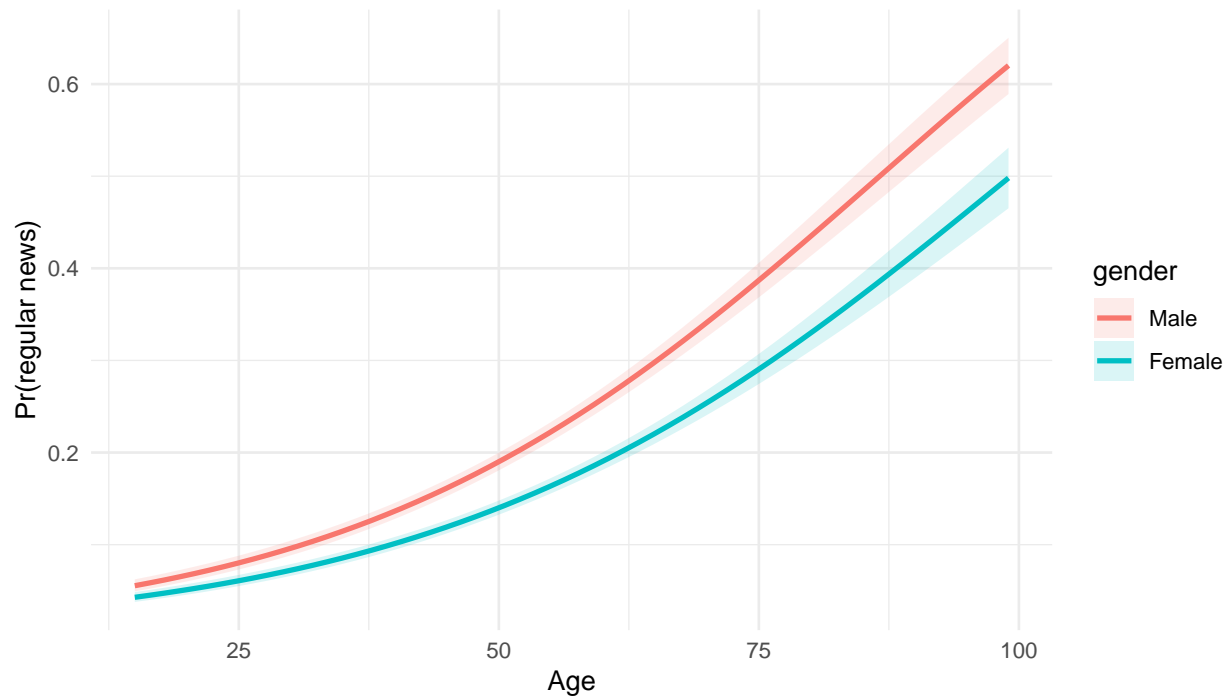
### 5.4 3. Binary $\times$ Continuous interaction (gender $\times$ age)

```
logit3 <- glm(news_regular ~ gender * agea + country + eduyrs, data = ess, family = binomial())
plot_gender_age <- ggplot(nd3, aes(x = agea, y = pr, color = gender)) +
  geom_line(size = 1) +
  labs(y = "Pr(regular news)", x = "Age") +
  theme_minimal()
```

#### 5.4.0.1 Code

#### 5.4.0.2 Output

```
## # A tibble: 7 x 7
##   term          estimate std.error statistic    p.value  conf.low  conf.high
##   <chr>          <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)    0.0109   0.114   -39.6    0        0.00868  0.0136
## 2 genderMale     1.26     0.113    2.03 4.22e- 2    1.01    1.57
## 3 agea           1.04     0.00141  26.2 1.61e-151  1.03    1.04
## 4 countryFR      0.733    0.0460   -6.76 1.34e- 11  0.669    0.802
## 5 countryGB      1.51     0.0370   11.2 3.74e- 29  1.41    1.63
## 6 eduyrs         1.03     0.00450    7.45 9.68e- 14  1.02    1.04
## 7 genderMale:agea 1.00     0.00192    1.41 1.58e- 1    0.999    1.01
```



Interpretation: Diverging ribbons indicate age effects differ by gender; if both ribbons rise similarly, the interaction is weak.

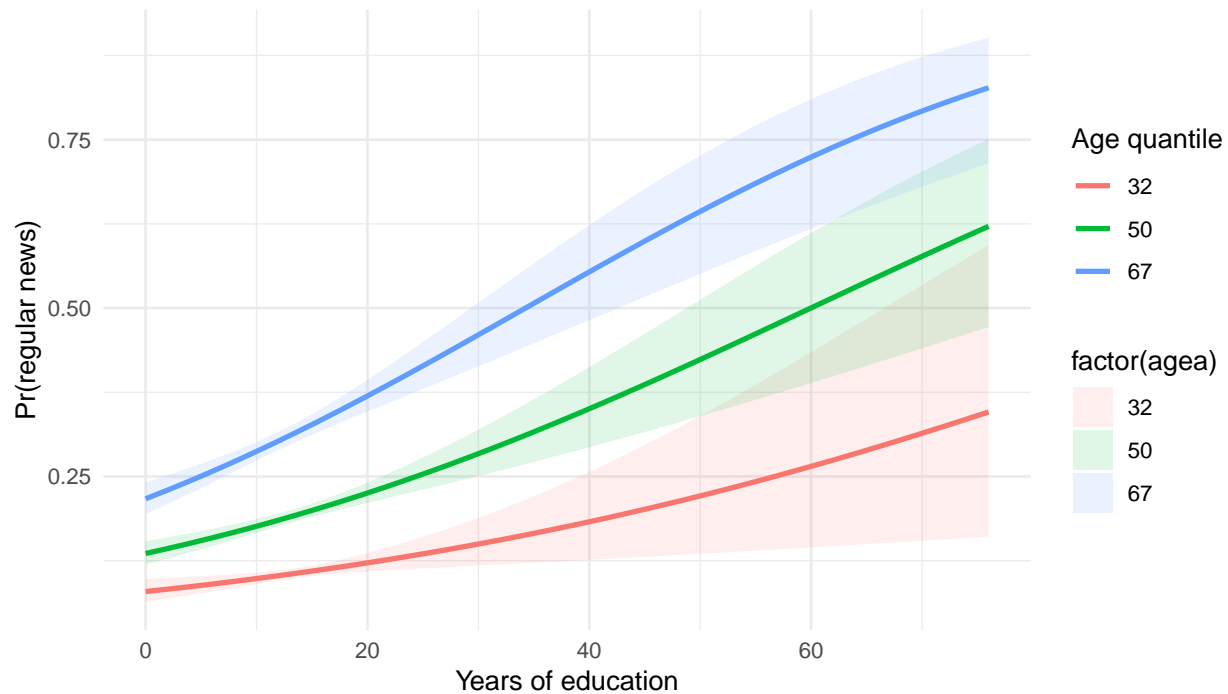
## 5.5 4. Continuous $\times$ Continuous interaction (age $\times$ education)

```
logit4 <- glm(news_regular ~ agea * eduyrs + gender + country, data = ess, family = binomial())
plot_age_edu <- ggplot(nd4, aes(x = eduyrs, y = pr, color = factor(agea))) +
  geom_line(size = 1) +
  labs(y = "Pr(regular news)", x = "Years of education", color = "Age quantile") +
  theme_minimal()
```

### 5.5.0.1 Code

### 5.5.0.2 Output

```
## # A tibble: 7 x 7
##   term          estimate std.error statistic  p.value  conf.low  conf.high
##   <chr>          <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)    0.0133   0.223    -19.4  1.12e-83  0.00860  0.0206
## 2 agea           1.03     0.00353   9.47  2.80e-21  1.03     1.04
## 3 eduyrs         1.01     0.0162    0.707  4.80e- 1  0.980    1.04
## 4 genderMale     1.46     0.0334   11.4   6.41e-30  1.37     1.56
## 5 countryFR      0.737    0.0461   -6.63  3.29e-11  0.673    0.806
## 6 countryGB      1.52     0.0370   11.2   2.53e-29  1.41     1.63
## 7 agea:eduyrs    1.00     0.000270  1.44  1.50e- 1  1.00     1.00
```



Interpretation: Steeper lines for higher age quantiles would mean education matters more (or less) for older respondents; overlap implies limited moderation.

## 5.6 5. Three-way interaction (gender $\times$ age $\times$ country)

```
logit5 <- glm(news_regular ~ gender * agea * country + eduysr, data = ess, family = binomial())
plot_threeway <- ggplot(nd5, aes(x = agea, y = pr, color = gender)) +
  geom_line() +
  facet_wrap(~ country) +
  labs(y = "Pr(regular news)", x = "Age") +
  theme_minimal()
```

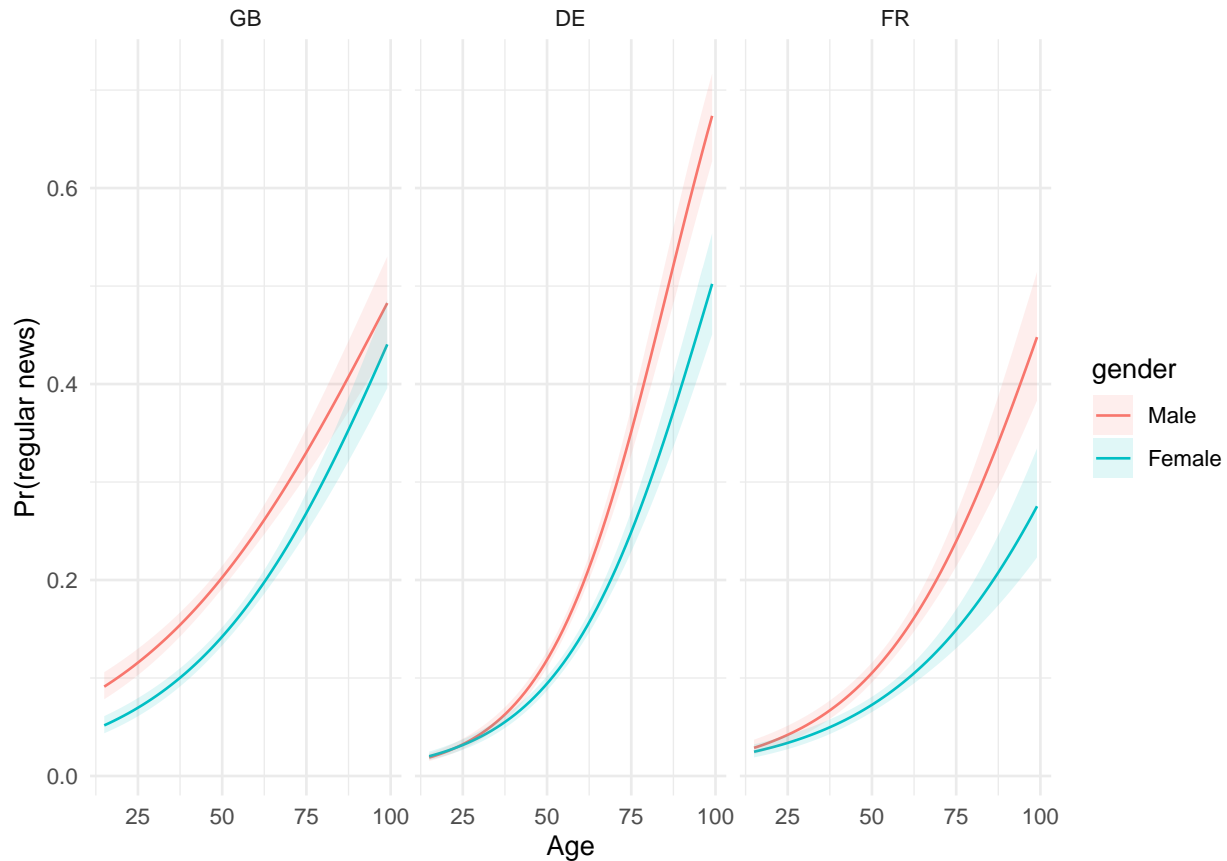
### 5.6.0.1 Code

### 5.6.0.2 Output

```
## # A tibble: 13 x 7
```

##	term	estimate	std.error	statistic	p.value	conf.low	conf.high
##	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
##	1 (Intercept)	0.00660	0.164	-30.7	2.20e-206	0.00477	0.00907
##	2 genderMale	0.809	0.203	-1.04	2.97e- 1	0.543	1.20
##	3 agea	1.05	0.00240	19.3	6.67e- 83	1.04	1.05
##	4 countryFR	1.52	0.232	1.82	6.91e- 2	0.965	2.40
##	5 countryGB	3.30	0.187	6.38	1.73e- 10	2.29	4.76
##	6 eduysr	1.03	0.00452	7.28	3.30e- 13	1.02	1.04
##	7 genderMale:agea	1.01	0.00336	2.79	5.33e- 3	1.00	1.02
##	8 genderMale:country~	1.29	0.324	0.793	4.28e- 1	0.685	2.44

```
## 9 genderMale:country~ 2.46      0.260      3.46 5.37e- 4 1.48      4.10
## 10 agea:countryFR      0.986      0.00387     -3.66 2.57e- 4 0.979      0.993
## 11 agea:countryGB      0.986      0.00311     -4.69 2.77e- 6 0.980      0.992
## 12 genderMale:agea:co~ 0.998      0.00542     -0.400 6.89e- 1 0.987      1.01
## 13 genderMale:agea:co~ 0.985      0.00438     -3.34 8.46e- 4 0.977      0.994
```



Interpretation: Scan each country facet—if ribbons separate widely, the age–gender pattern is country-specific; overlapping ribbons suggest similar patterns across countries.

## 5.7 6. Marginal effects and interpretation

```
# Average marginal effects for age within each country
ame_age <- marginaleffects(logit5, variables = "agea", by = "country")

# Marginal effect of being Female at age 25 vs age 65
me_female_age <- marginaleffects(logit5, variables = "gender", newdata = datagrid(agea = c(25, 65)))
```

- Prefer predicted probabilities and marginal effects over raw log-odds.
- Inspect separation or influential points with `performance::check_model(logit5)` if desired.

### 5.7.1 AMEs and focal contrasts

```

ame_table <- tibble(
  variable = c("agea (+1 year)", "gender (Female vs Male)"),
  AME = c(ame_age, ame_fem)
)

me_female_diff <- data.frame(agea = c(25,65), diff = pred_f - pred_m) # from fits chunk

```

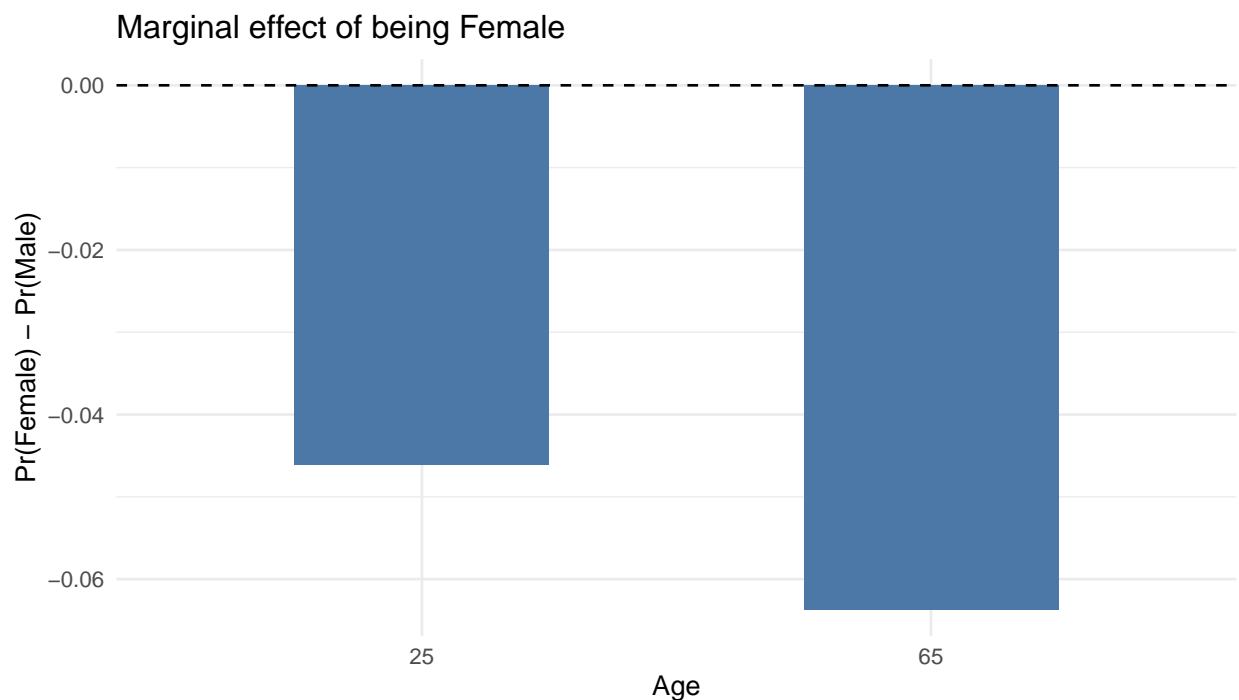
#### 5.7.1.1 Code

#### 5.7.1.2 Output

```

## # A tibble: 2 x 2
##   variable          AME
##   <chr>          <dbl>
## 1 agea (+1 year)  0.00451
## 2 gender (Female vs Male) 0.00237

```



Interpretation: The AME for age reports the average change in the probability of regular news use for a one-year increase in age. The bar chart shows how the female–male gap differs at ages 25 vs 65; positive bars mean higher probability for women at that age.

Country-level pooling choices (fixed vs multilevel) are expanded in the next chapter.

## 5.8 Problem set — Logistic regression practice

1. Recode the outcome as `news_daily = nwsptot >= 5` and re-estimate `logit3`. How do the marginal effects of age change when the bar for “regular” consumption is higher?

2. Add `urban` (1/2 = urban, 3-5 = non-urban) as a predictor and interact it with `country`. Which country shows the largest urban-rural gap in news readership?
3. Compare `logit4` and `logit5` using AIC and pseudo  $R^2$  (`pscl::pR2`). Which balance of complexity and fit seems reasonable for classroom examples?
4. For one model, translate results into plain language for a non-technical audience: choose two profiles (e.g., 30-year-old woman in GB vs 60-year-old man in FR) and report predicted probabilities.

These exercises deliver a full set of interaction types in a nonlinear setting, ready to complement your lecture slides.

## 6 Country effects — fixed effects and multilevel modeling

Nested survey data call for explicit country handling. This chapter shows two approaches:

- **Country fixed effects (dummies):** absorbs unobserved, time-invariant differences between GB, DE, and FR.
- **Multilevel (random-intercept) logistic regression:** partially pools country effects, reducing noise for small samples and enabling variance decomposition.

```
library(dplyr)
library(ggplot2)
library(broom)
library(broom.mixed)
library(lme4)
library(tidyr)

source("R/clean_ess.R")

ess <- clean_ess()
```

### 6.1 1. Country fixed effects (logit with dummies)

- **Model:** `news_regular ~ agea + gender + eduyrs + country`
- **Interpretation:** Country coefficients capture average gaps relative to the reference country (DE, alphabetically first).

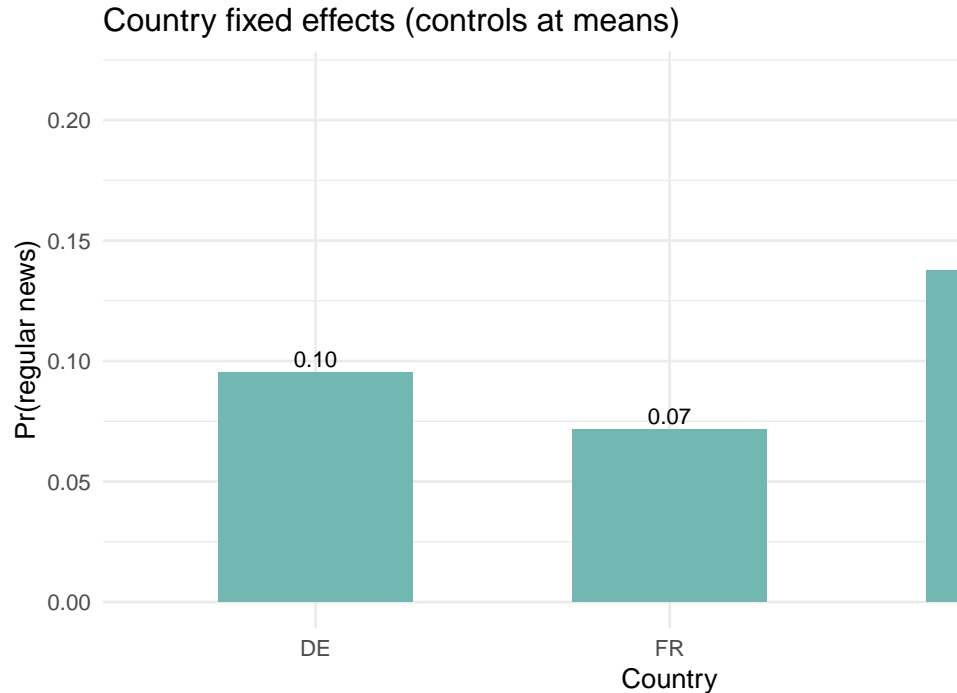
```
fe_logit <- glm(news_regular ~ agea + gender + eduyrs + country,
               data = ess, family = binomial())
fe_tidy <- broom::tidy(fe_logit, exponentiate = TRUE, conf.int = TRUE)
```

#### 6.1.0.1 Odds ratios

##### 6.1.0.1.1 Table

```
## # A tibble: 6 x 7
##   term          estimate std.error statistic  p.value conf.low conf.high
##   <chr>          <dbl>     <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)  0.00997  0.0971   -47.5    0        0.00824  0.0121
## 2 agea         1.04     0.000998  38.4    0        1.04     1.04
```

## 3	genderMale	1.47	0.0334	11.5	1.57e-30	1.37	1.57
## 4	eduyrs	1.03	0.00449	7.55	4.36e-14	1.03	1.04
## 5	countryFR	0.733	0.0460	-6.75	1.49e-11	0.670	0.802
## 6	countryGB	1.51	0.0370	11.2	3.56e-29	1.41	1.63



#### 6.1.0.1.2 Country contrasts (plot)

Takeaway: Fixed effects wipe out between-country bias but treat each country independently—estimates can be noisy if a country has few cases.

## 6.2 2. Multilevel logistic model (random intercepts by country)

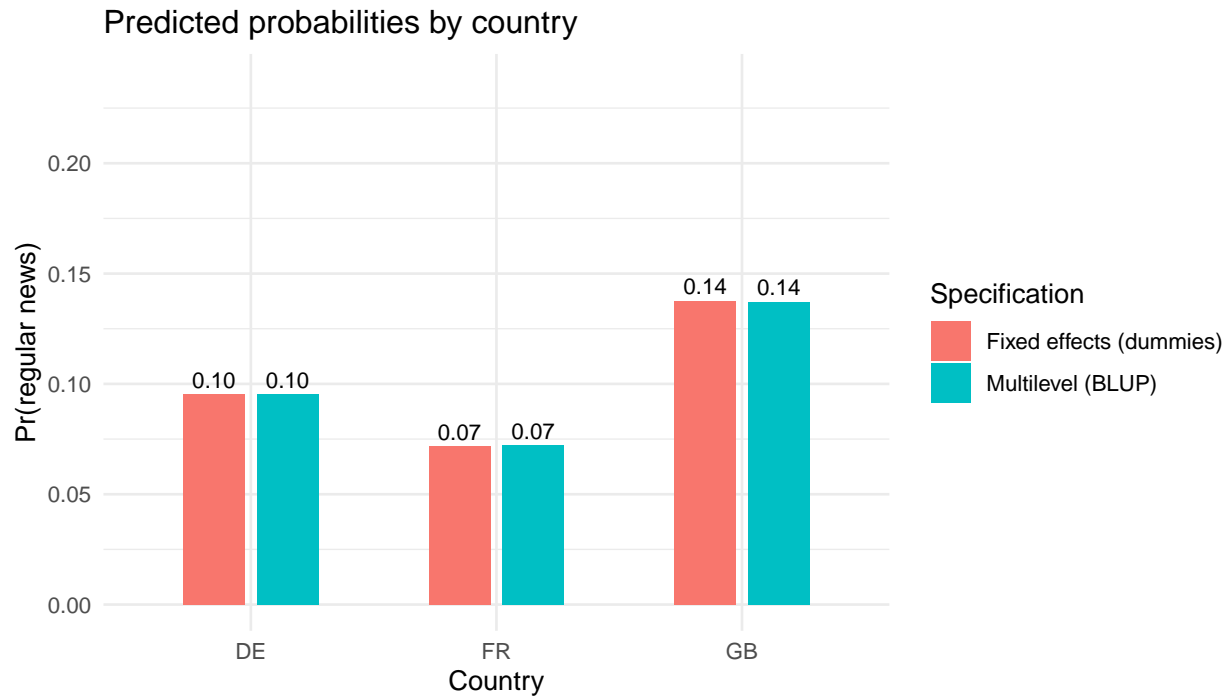
- **Model:** `news_regular ~ agea + gender + eduyrs + (1 | country)`
- **Why:** Partial pooling shrinks extreme country estimates toward the grand mean, improving out-of-sample stability.

```
ml_logit <- glmer(news_regular ~ agea + gender + eduyrs + (1 | country),
  data = ess, family = binomial(),
  control = glmerControl(optimizer = "bobyqa"))

var_u0 <- as.numeric(VarCorr(ml_logit)$country)
icc <- var_u0 / (var_u0 + pi^2 / 3)
```

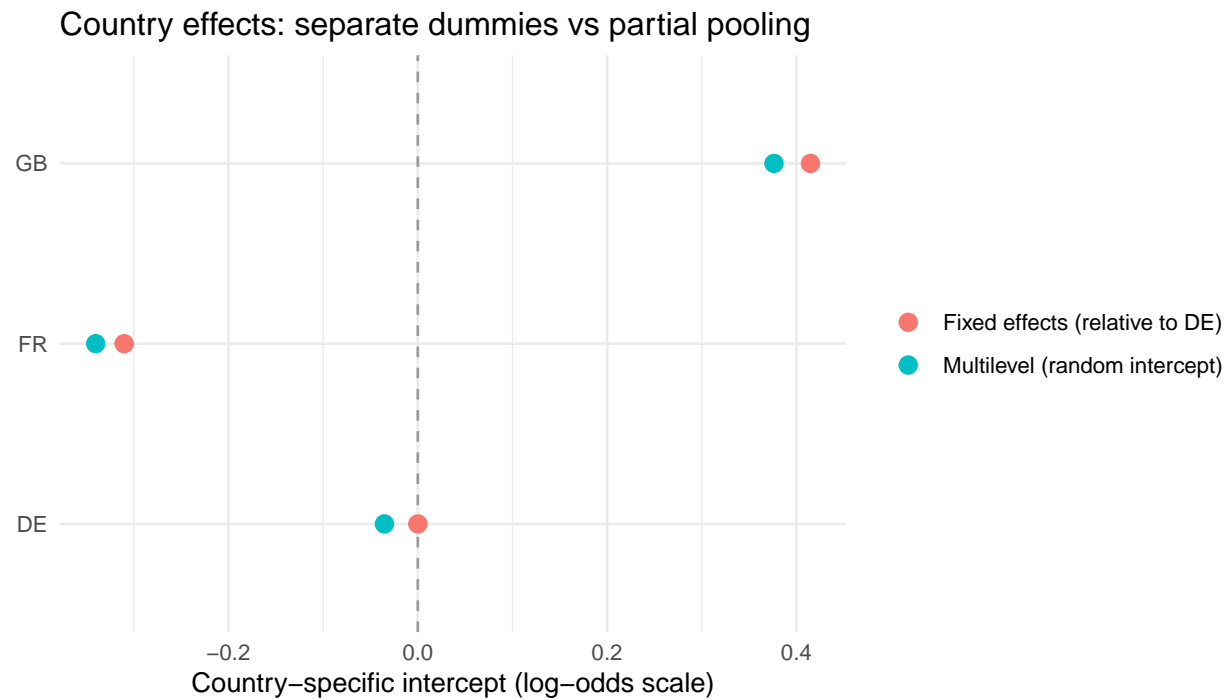
Intraclass correlation (ICC): 2.58% of the variance in the log-odds of regular news use is at the country level.

### 6.2.1 Country predictions: fixed vs multilevel



Interpretation: Multilevel estimates are closer together because extreme country effects are shrunk toward the overall mean.

### 6.2.2 Visualizing country intercepts (shrinkage)





Reading the plot: Dots further from 0 indicate larger country-specific deviations. The multilevel model shrinks them toward zero, while fixed effects leave them unchanged.

### 6.3 3. Random slopes for gender (optional extension)

With more countries, we could allow the gender gap to vary by country:

```
ml_logit_gender <- glmer(  
  news_regular ~ agea + gender + eduyrs + (1 + gender | country),  
  data = ess, family = binomial(),  
  control = glmerControl(optimizer = "bobyqa")  
)
```

In the current three-country sample this model may be over-parameterized; the random-intercept specification above is the stable classroom default.

### 6.4 Practice prompts

1. Add an **urban** fixed effect to both models. Do city–rural gaps widen or narrow once country pooling is applied?
2. Refit the multilevel model with **news\_regular** defined as daily readership (**nwsptot** >= 5). How does the ICC change?
3. Replace **agea** with a spline (**splines::ns(agea, df = 3)**) inside both models and compare the resulting age profiles by country.

## 7 Survey weighting — why and how

Survey data are collected with unequal selection probabilities. Inference should reflect the design to avoid biased point estimates and standard errors.

- **Notation:** Let  $w_i = 1/\pi_i$  be the design (inverse-probability) weight. For a finite population mean  $\bar{Y} = \sum_i w_i Y_i / \sum_i w_i$ . For regression, weighted likelihoods re-scale each case by  $w_i$ .
- **ESS fields used:** **pweight** (post-stratification weight), **psu** (primary sampling unit), **stratum** (strata). We set `options(survey.lonely.psu = "adjust")` to stabilize single-PSU strata.

### 7.0.1 What these design variables mean (plain language)

- **pweight (post-stratification weight):** Adjusts for unequal inclusion probabilities *and* aligns the achieved sample with known population margins (e.g., age × gender × region). Large **pweight** values upweight under-represented respondents; small values downweight over-represented ones.
- **psu (primary sampling unit):** The first cluster stage of selection (e.g., municipalities or postcode sectors). Respondents inside the same PSU share fieldwork and selection features, so their responses are correlated.
- **stratum (strata):** Mutually exclusive groups within which PSUs were sampled (e.g., by region × urbanicity). Stratification improves precision; variance estimation must respect it.
- **Design degrees of freedom:** With clustering and stratification, the effective df are closer to the number of PSUs minus strata, not the raw respondent count—hence the importance of design-aware SEs.

```

library(dplyr)
library(ggplot2)
library(tidyr)
library(survey)
library(broom)

# Synthetic microdata to illustrate weighting without depending on raw ESS file quirks
set.seed(42)
n_psu <- 120
n_by_psu <- sample(8:18, n_psu, replace = TRUE)

demo_psu <- tibble(
  psu = 1:n_psu,
  stratum = sample(1:20, n_psu, replace = TRUE),
  country = sample(c("GB", "DE", "FR"), n_psu, replace = TRUE, prob = c(.4, .35, .25)),
  pweight = runif(n_psu, 0.5, 3),
  n = n_by_psu
)

ess_w <- demo_psu |>
  uncount(n) |>
  mutate(
    agea = round(rnorm(n(), 50, 15)),
    gender = sample(c("Male", "Female"), n(), replace = TRUE),
    # Generate outcome with country- and age-dependent probability
    linpred = -1 + 0.6 * (country == "GB") + 0.3 * (country == "FR") +
      0.01 * (agea - 50) + 0.4 * (gender == "Female"),
    news_regular = rbinom(n(), 1, plogis(linpred))
  ) |>
  select(psu, stratum, country, pweight, agea, gender, news_regular) |>
  mutate(country = factor(country),
    gender = factor(gender))

options(survey.lonely.psu = "adjust")
des <- svydesign(ids = ~psu, strata = ~stratum, weights = ~pweight,
  data = ess_w, nest = TRUE)

svy_n <- nrow(des$variables)

```

## 7.1 1. Weighted descriptive estimates

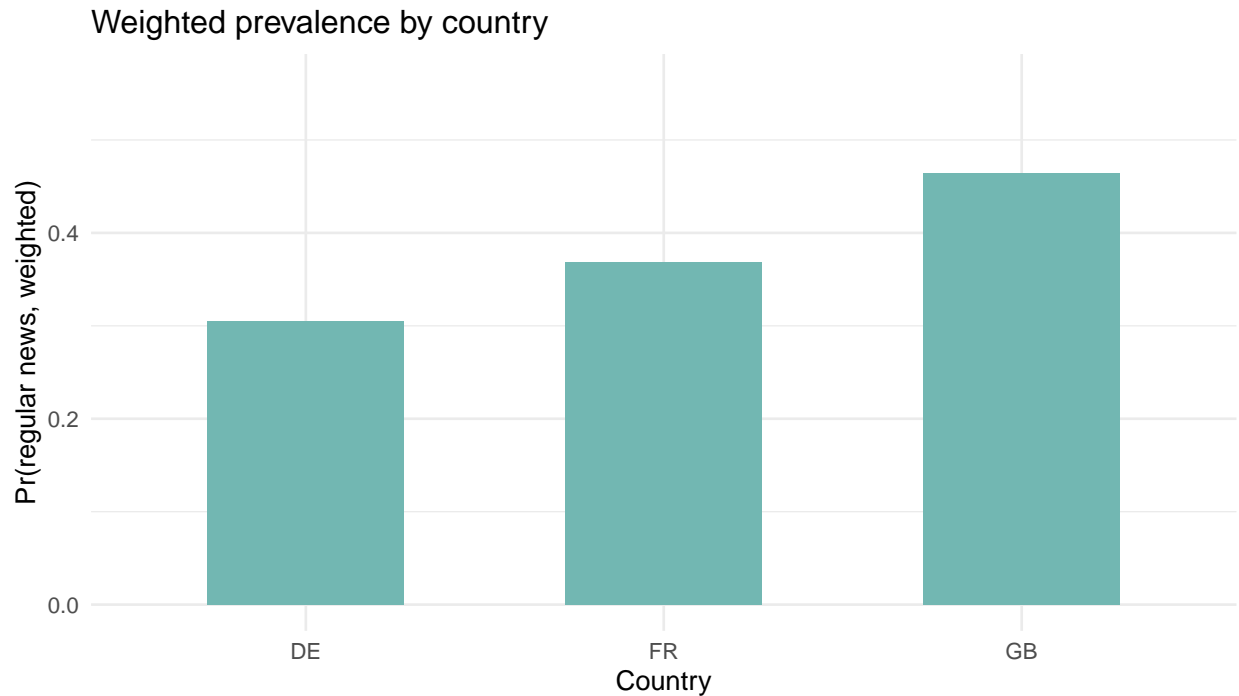
Sample used after complete-case filtering: 1556 respondents.

```
svymean(~news_regular, des)
```

```
##              mean      SE
## news_regular 0.38522 0.0153
```

- The point estimate is the **design-weighted mean**; SEs account for clustering and stratification.
- **Design effect** (`svymean(..., deff=TRUE)`) tells how much variance inflation comes from the design versus SRS.

### 7.1.1 Country-weighted proportions



## 7.2 2. Weighted regression: linear probability and logit

```
lpm_w <- svyglm(news_regular ~ agea + gender + country,  
                design = des, family = gaussian())  
tidy(lpm_w)
```

```
## # A tibble: 5 x 5  
##   term      estimate std.error statistic    p.value  
##   <chr>      <dbl>    <dbl>    <dbl>    <dbl>  
## 1 (Intercept) 0.240    0.0493     4.86 0.00000453  
## 2 agea        0.00214  0.000781    2.74 0.00741  
## 3 genderMale -0.0875   0.0243    -3.60 0.000509  
## 4 countryFR   0.0679   0.0390     1.74 0.0851  
## 5 countryGB   0.155    0.0276     5.61 0.000000193
```

```
logit_w <- svyglm(news_regular ~ agea + gender + country,  
                  design = des, family = quasibinomial())  
tidy(logit_w, exponentiate = TRUE)
```

```
## # A tibble: 5 x 5  
##   term      estimate std.error statistic    p.value  
##   <chr>      <dbl>    <dbl>    <dbl>    <dbl>  
## 1 (Intercept) 0.327    0.221    -5.06 0.00000204  
## 2 agea        1.01    0.00346    2.69 0.00833  
## 3 genderMale  0.683    0.105    -3.64 0.000446
```

## 4	countryFR	1.36	0.174	1.76	0.0815
## 5	countryGB	1.95	0.122	5.48	0.000000335

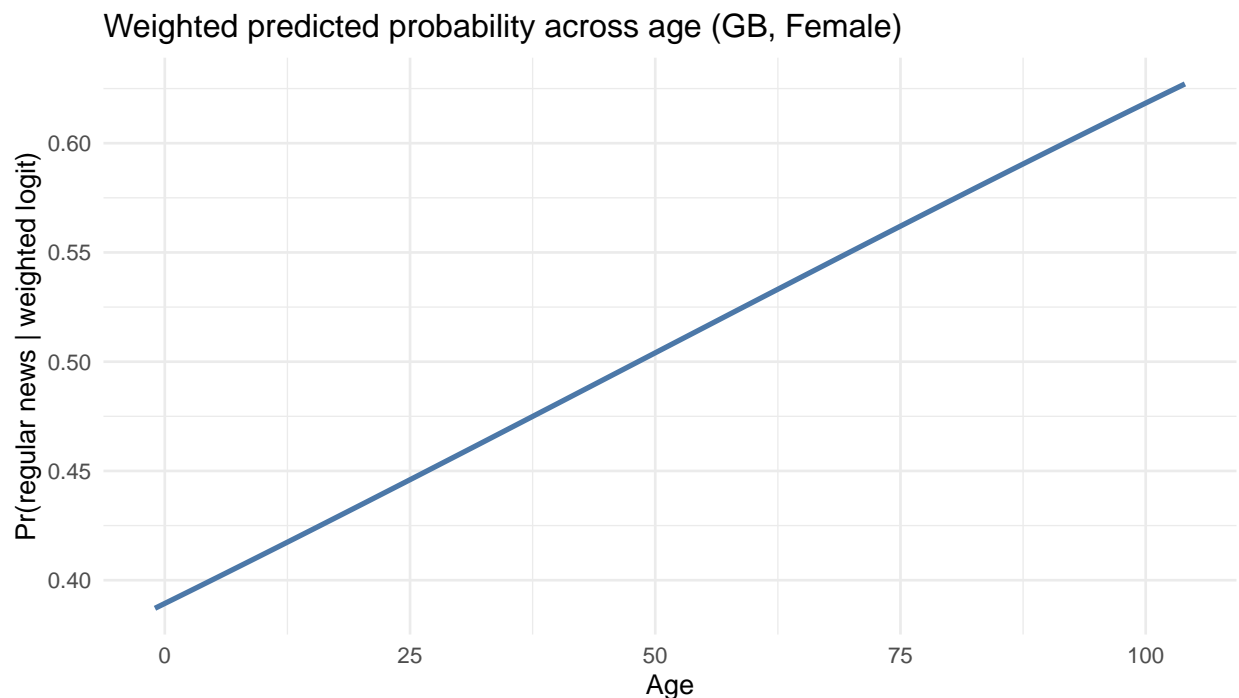
Comparison points for students:

- **Weights + clustering:** `svyglm` gives design-consistent SEs; plain `glm` does not.
- **Quasibinomial** keeps logit link but uses robust variance; estimates mirror survey-weighted MLE when weights are scaled.
- **LPM vs logit under weights:** LPM slopes stay probability-difference interpretation; logit ORs remain multiplicative.

### 7.3 3. When to weight (practical guidance)

- Use design/post-strat weights when estimating **population levels** (means, totals, prevalence) or effects that might shift with differential selection.
- In randomized experiments or when modeling **causal effects with ignorable sampling**, weights may be optional; still cluster-robust SEs matter.
- If the research question is **sample-only prediction**, weights can be skipped, but report that scope-of-inference is limited.

### 7.4 4. Small worked example: weighted marginal effect of age



### 7.5 Practice prompts

1. Recompute weighted country gaps using `anweight` instead of `pweight`. How do the estimates move?
2. Add education to the weighted logit. Do ORs shift relative to the unweighted model in Chapter 5?
3. Estimate a design effect for `news_regular` and discuss how many “effective” observations the design corresponds to.

## 8 Appendix — Data and build notes

### 8.1 ESS subset

- File: `ess.csv`
- Countries: GB, DE, FR
- For variable descriptions, open the bundled HTML codebook: `ESS1e06_...subset codebook.html` in your browser.
- Common missing-value codes: 66/77/88/99 or blank strings. All chapters use `na_if()` to drop them before analysis.

### 8.2 Extending the material

- Swap dependent variables: `pplhlp` (people helpful) or `pplfair` (people fair) can replace `ppltrst` without changing structure.
- Add country-level covariates by merging lookups (e.g., GDP or media freedom indices) before running multilevel models.
- Convert interaction plots to `ggplotly` for live demos if desired.