

Statistics for Linguists with R – a SIGIL course

Unit 8: Non-Randomness of Corpus Data & Generalised Linear Models

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<http://purl.org/stefan.evert/SIGIL>

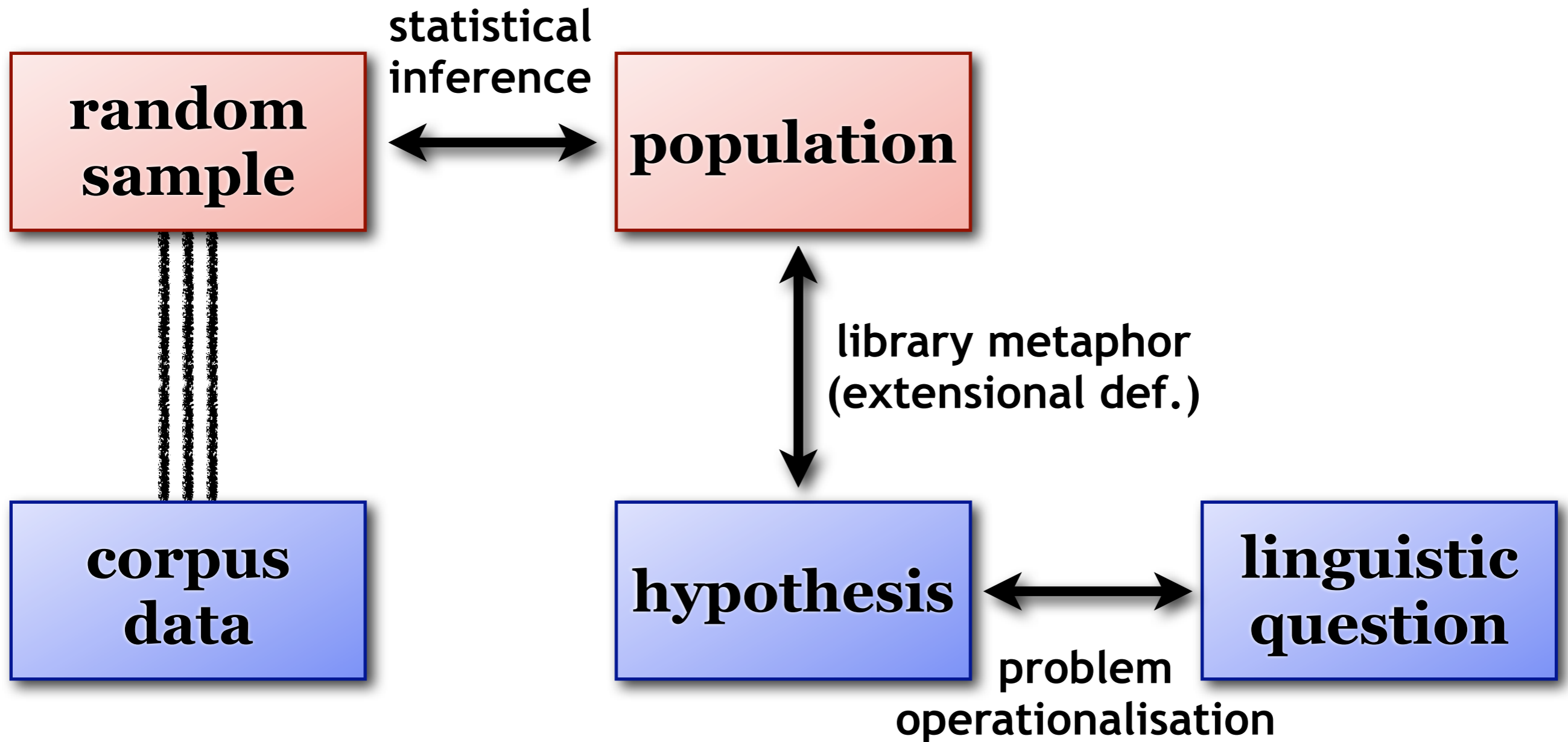
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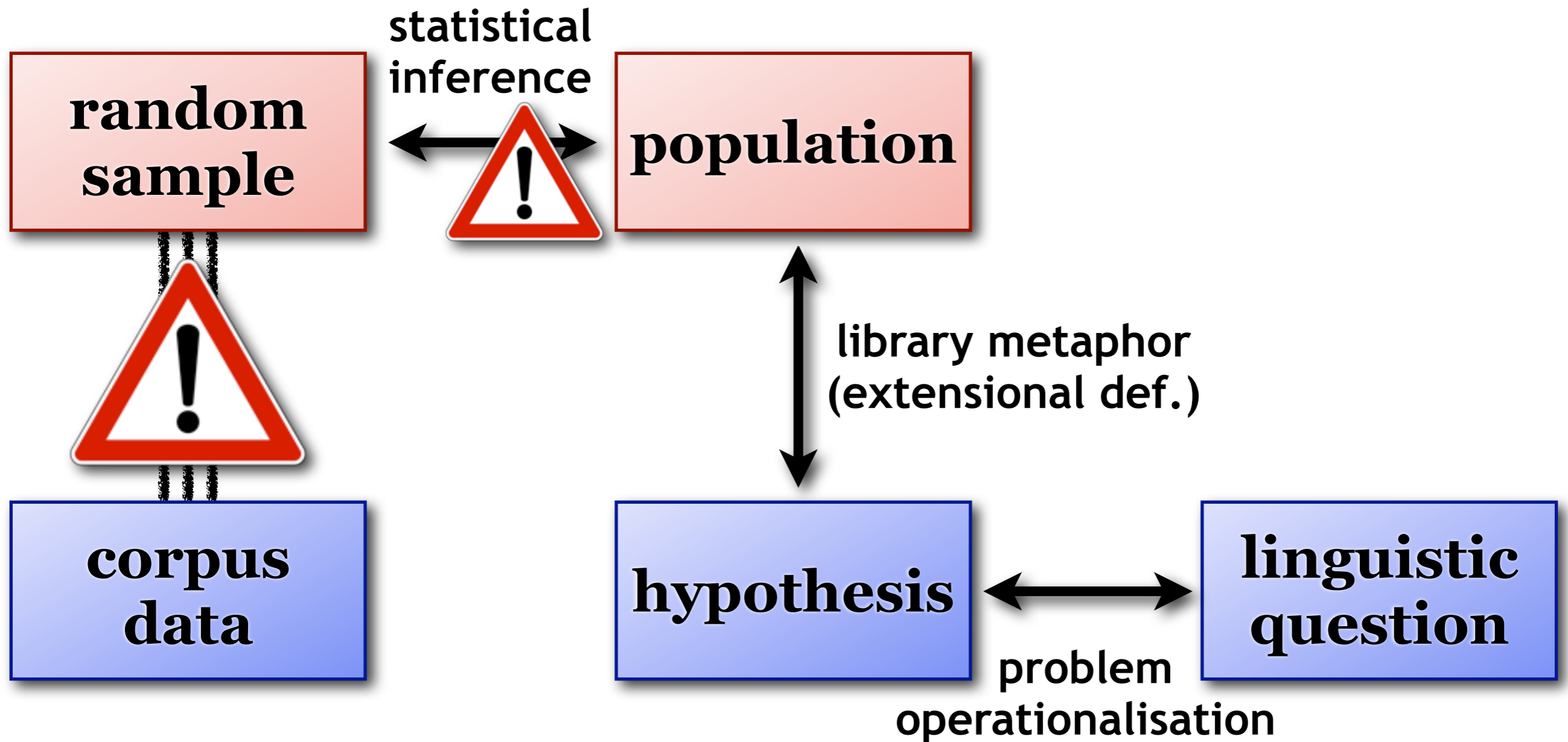


Introduction & Reminder

Problems with statistical inference



Problems with statistical inference



Mathematical problems: Significance

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 - especially highly skewed tables in collocation extraction
 - G^2 overestimates significance for small samples (well-known in statistics, e.g. Agresti 2002)
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 - G^2 overestimates significance for small samples (well-known in statistics, e.g. Agresti 2002)
 - e.g. manual samples of 100–500 items (as in our examples)
 - often ignored because of its success in computational linguistics
 - Fisher is conservative & computationally expensive
 - also numerical problems, e.g. in R version 1.x 

Mathematical problems:

Effect size

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- ◆ Effect size for frequency comparison
 - not clear which measure of effect size is appropriate
 - e.g. **difference** of proportions, **relative risk** (ratio of proportions), **odds ratio**, logarithmic odds ratio, normalised χ^2 , ...

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 - not clear which measure of effect size is appropriate
 - e.g. **difference** of proportions, **relative risk** (ratio of proportions), **odds ratio**, logarithmic odds ratio, normalised χ^2 , ...
- ◆ Confidence interval estimation
 - accurate & efficient estimation of confidence intervals for effect size is often very difficult
 - exact confidence intervals only available for odds ratio

Mathematical problems: Multiple hypothesis tests

- ◆ Each individual hypothesis test controls risk of type I error ... but if you carry out thousands of tests, some of them *have* to be false rejections
 - recommended reading: *Why most published research findings are false* (Ioannidis 2005)
 - a monkeys-with-typewriters scenario

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*if applied to a single candidate selected *a priori**
 - but usually candidates selected *a posteriori* from data
→ many “unreported” tests for candidates with $f = 0$!
 - large number of such word pairs according to **Zipf's law** results in substantial number of type I errors
 - can be quantified with LNRE models (Evert 2004),
cf. Unit 5 on *word frequency distributions with zipfR*



Why a corpus isn't a random sample

Corpora

Corpora

- ◆ Theoretical sampling procedure is impractical
 - it would be very tedious if you had to take a random sample from a library, especially a hypothetical one, every time you want to test some hypothesis
- ◆ Use pre-compiled sample: a **corpus**

Corpora

- ◆ Theoretical sampling procedure is impractical
 - it would be very tedious if you had to take a random sample from a library, especially a hypothetical one, every time you want to test some hypothesis
- ◆ Use pre-compiled sample: a **corpus**
 - but this is not a random sample of tokens!
 - would be prohibitively expensive to collect 10 million VPs for a BNC-sized sample at random
 - other studies will need tokens of different granularity (words, word pairs, sentences, even full texts)

The Brown corpus

- ◆ First large-scale electronic corpus
 - compiled in 1964 at Brown University (RI)
- ◆ 500 samples of approx. 2,000 words each
 - sampled from edited AmE published in 1961
 - from 15 domains (imaginative & informative prose)
 - manually entered on punch cards

The British National Corpus

- ◆ 100 M words of modern British English
 - compiled mainly for lexicographic purposes: Brown-type corpora (such as LOB) are too small
 - both written (90%) and spoken (10%) English
 - XML edition (version 3) published in 2007
- ◆ 4048 samples from 25 to 428,300 words
 - 13 documents < 100 words, 51 > 100,000 words
 - some documents are collections (e.g. e-mail messages)
 - rich metadata available for each document

Unit of sampling

- ◆ Key problem: **unit of sampling** (text or fragment) \neq **unit of measurement** (e.g. VP)
 - recall sampling procedure in library metaphor ...

Unit of sampling



Unit of sampling

- ◆ Random sampling in the library metaphor
 - walk to a random shelf ...
 - ... select a random book ...
 - ... open it on a random page ...
 - ... and pick a random sentence from the page
 - ➔ repeat n times for sample size n

Unit of sampling

- ◆ Random sampling in the library metaphor
 - walk to a random shelf ...
 - ... select a random book ...
 - ... open it on a random page ...
 - ... and pick a random sentence from the page
 - ➔ repeat n times for **sample size n**
- ◆ Corpus = random sample of books, not sentences!
 - we should only use 1 sentence from each book
 - ➔ sample size: $n=500$ (Brown) or $n=4048$ (BNC)

Pooling data

- ◆ In order to obtain larger samples, researchers usually **pool** all data from a corpus
 - i.e. they include all sentences from each book
- ◆ Do you see why this is wrong?

Pooling data

- ◆ Books aren't random samples themselves
 - each book contains relatively homogeneous material
 - but much larger differences *between* books
- ◆ Therefore, the pooled data do not form a random sample from the library
 - for each randomly selected sentence, we co-select a substantial amount of very similar material
- ◆ Consequence: sampling variation increased

Pooling data

Pooling data

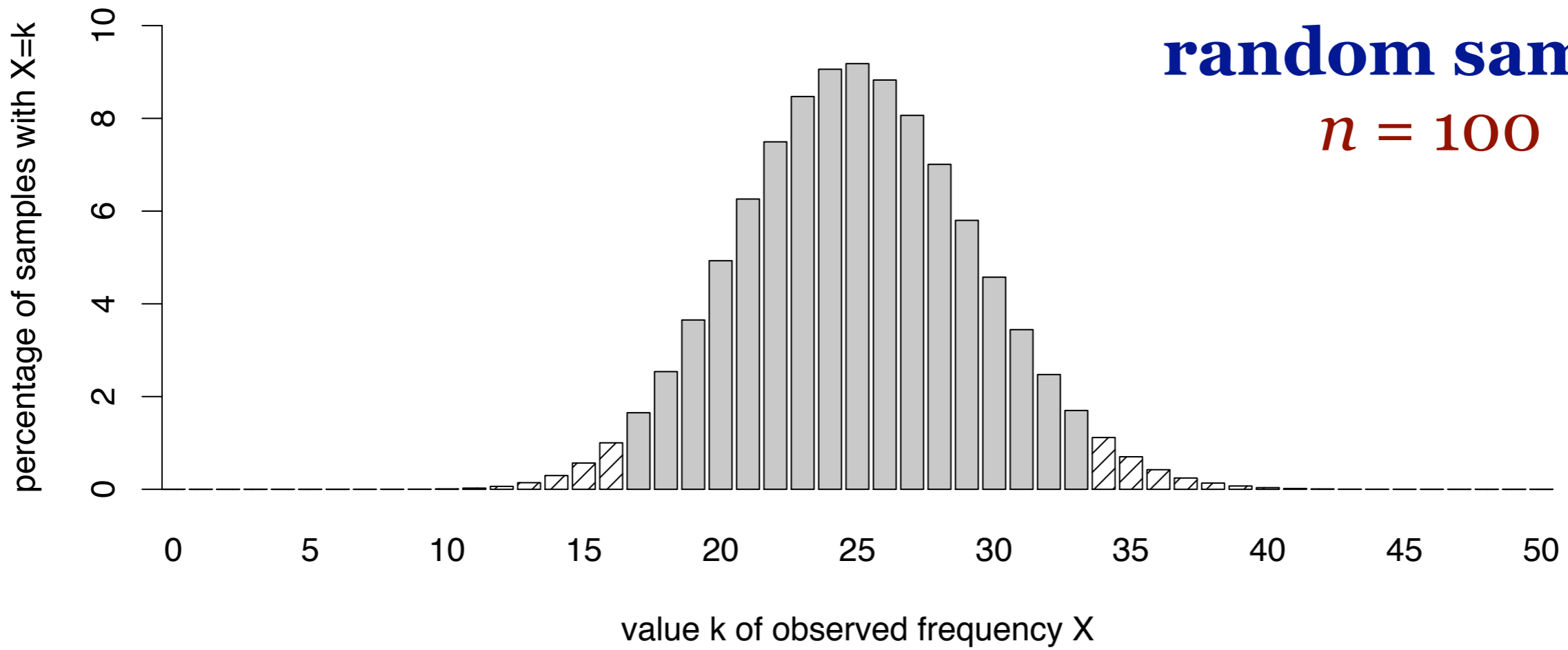
- ◆ Let us illustrate this with a simple example ...
 - assume library with two sections of equal size
 - e.g. spoken and written language in a corpus
 - population proportions are 10% vs. 40%
 - overall proportion of $\pi = 25\%$ in the library
 - this is the null hypothesis H_0 that we will be testing

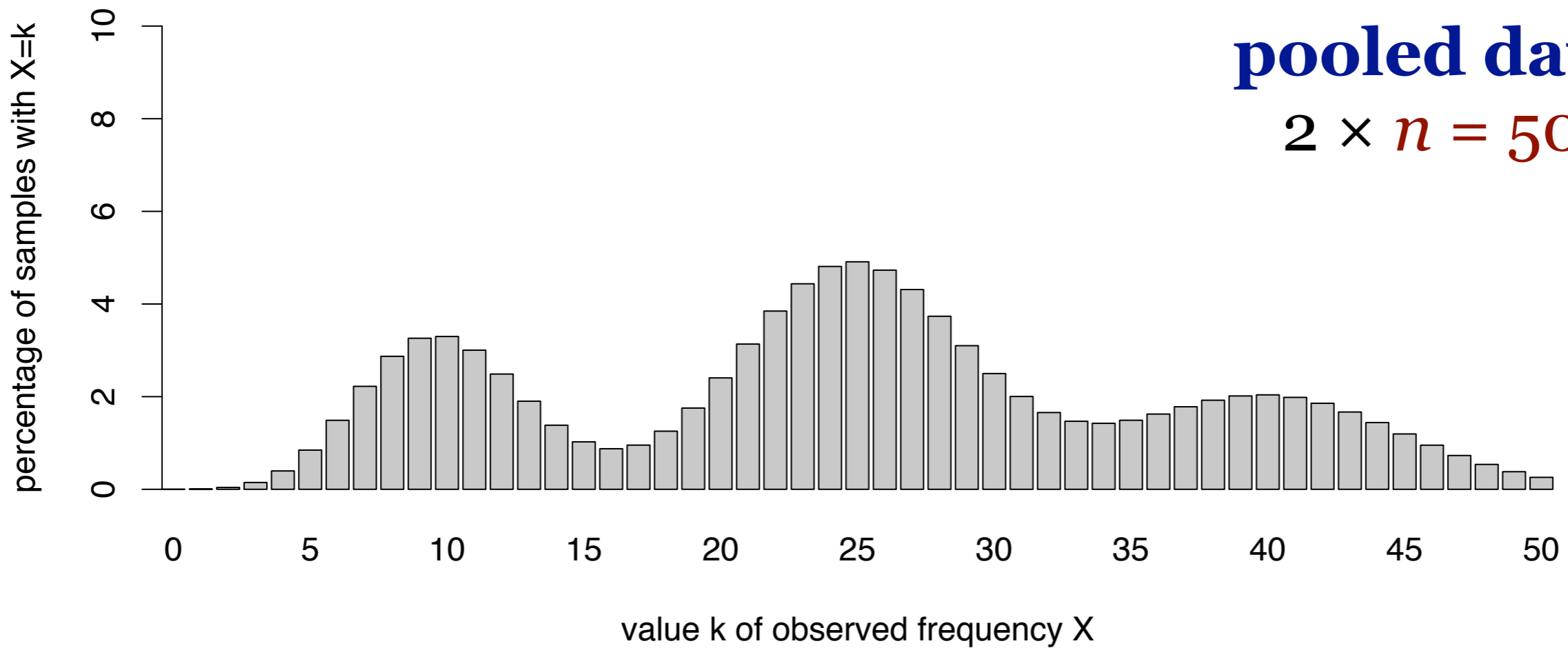
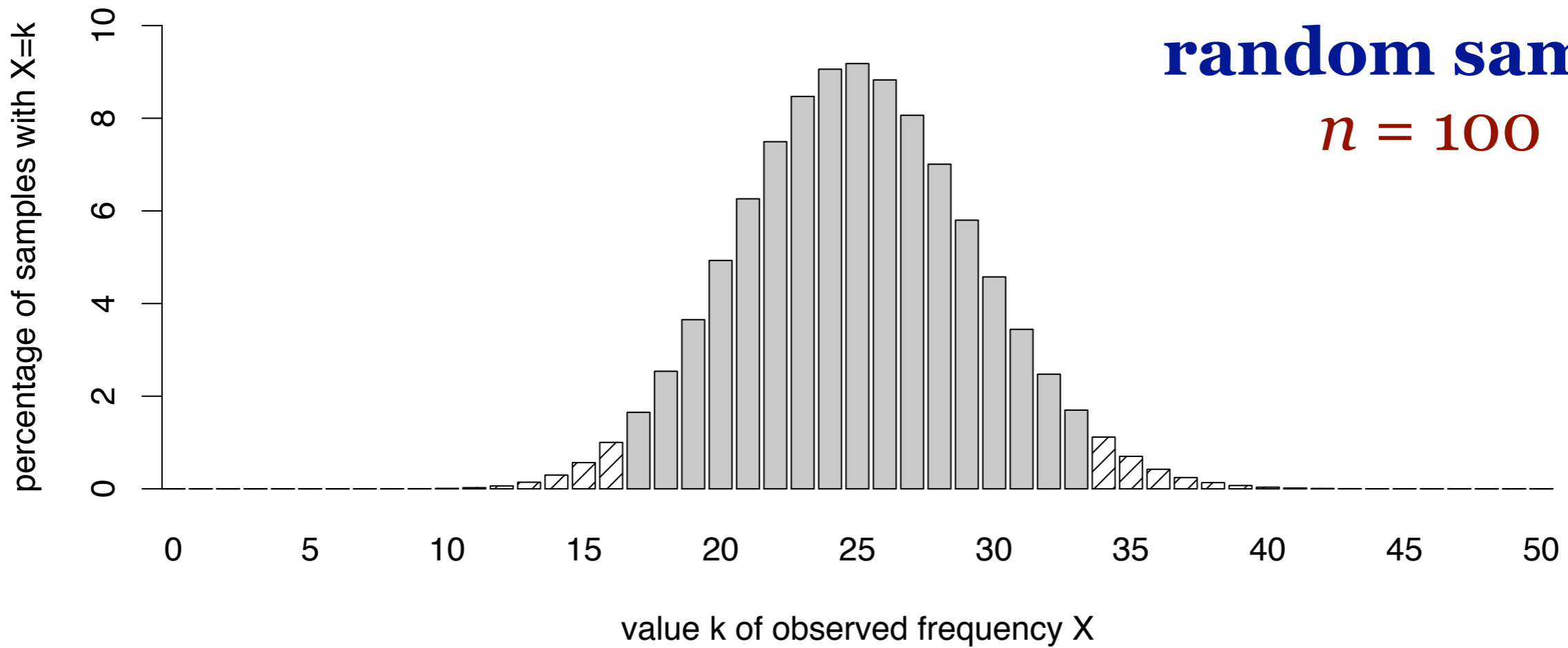
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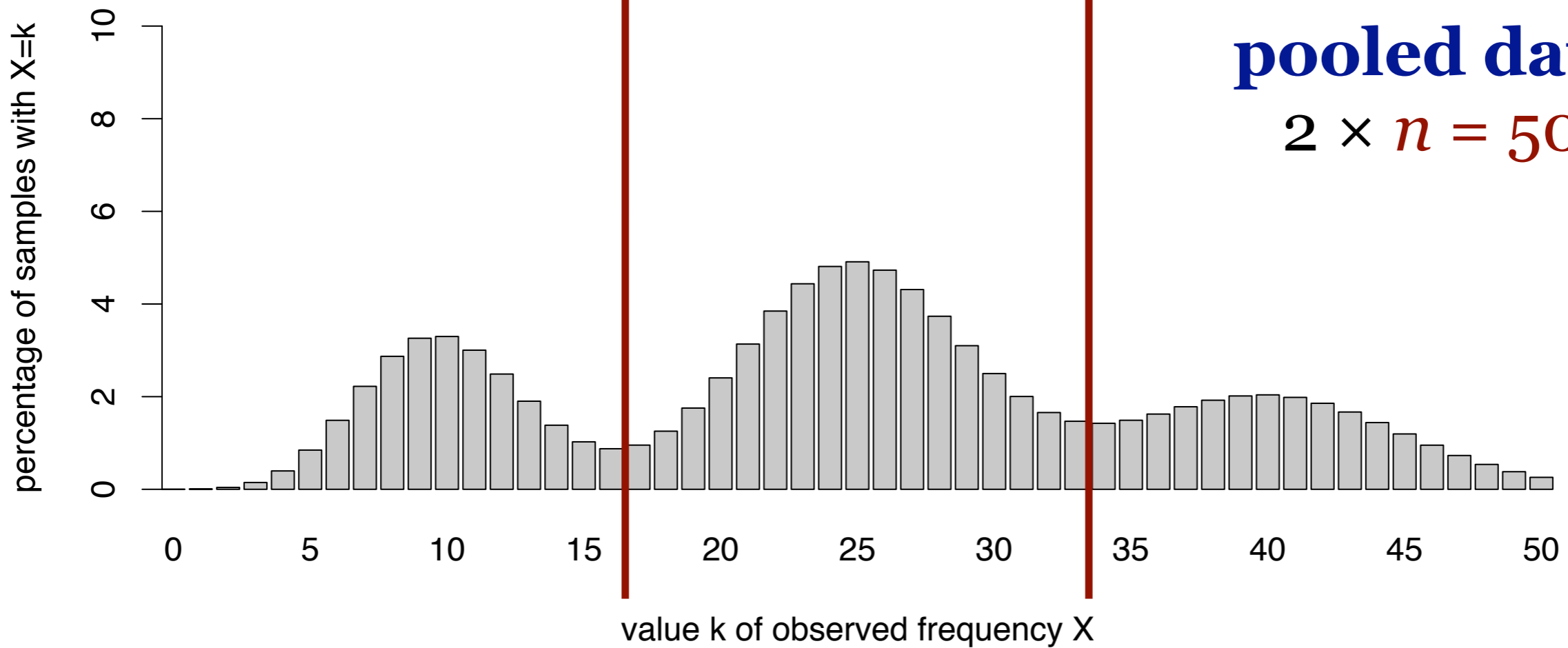
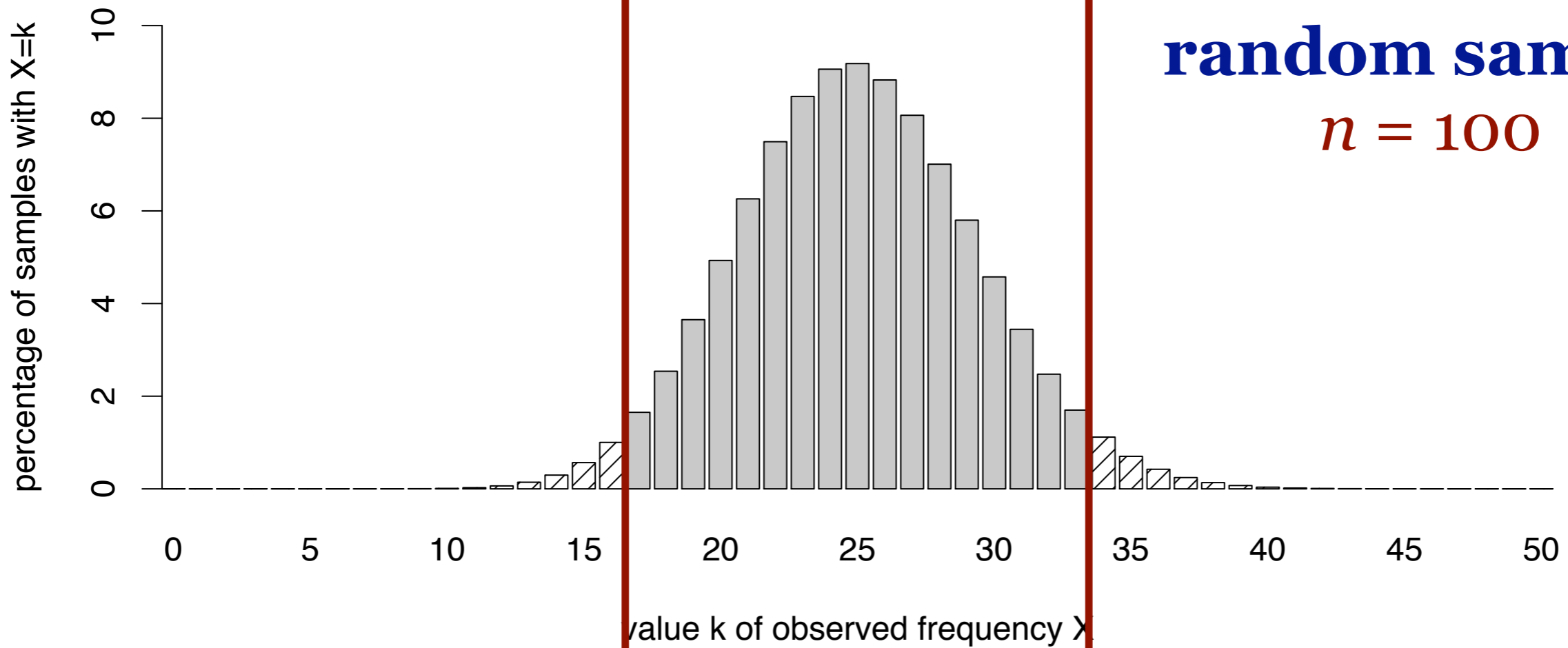
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 - this is the null hypothesis H_0 that we will be testing
- ◆ Compare sampling variation for
 - random sample of 100 tokens from the library
 - two randomly selected books of 50 tokens each
 - book is assumed to be a random sample from its section

random sample

$n = 100$







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 - 117 (!) occurrences in BNC, all in file HWX
 - very difficult to detect automatically
- ◆ Even worse for newspapers & Web corpora
 - see Evert (2004) for examples



Measuring non-randomness

A sample of random samples is a random sample

- ◆ Larger unit of sampling is not the original cause of non-randomness
 - if each text in a corpus is a genuinely random sample from the same population, then the pooled data also form a random sample
 - we can illustrate this with a thought experiment

The random library



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 - repeat until the heap of sentences is gone
- ➔ library of **random samples**
- ◆ Pooled data from 2 (or more) boxes form a perfectly **random sample** of sentences from the original library!



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 - discrepancy between unit of sampling and unit of measurement only leads to non-randomness if the sampling units (i.e. the corpus texts) are not random samples themselves (from same population)
 - with respect to specific phenomenon of interest

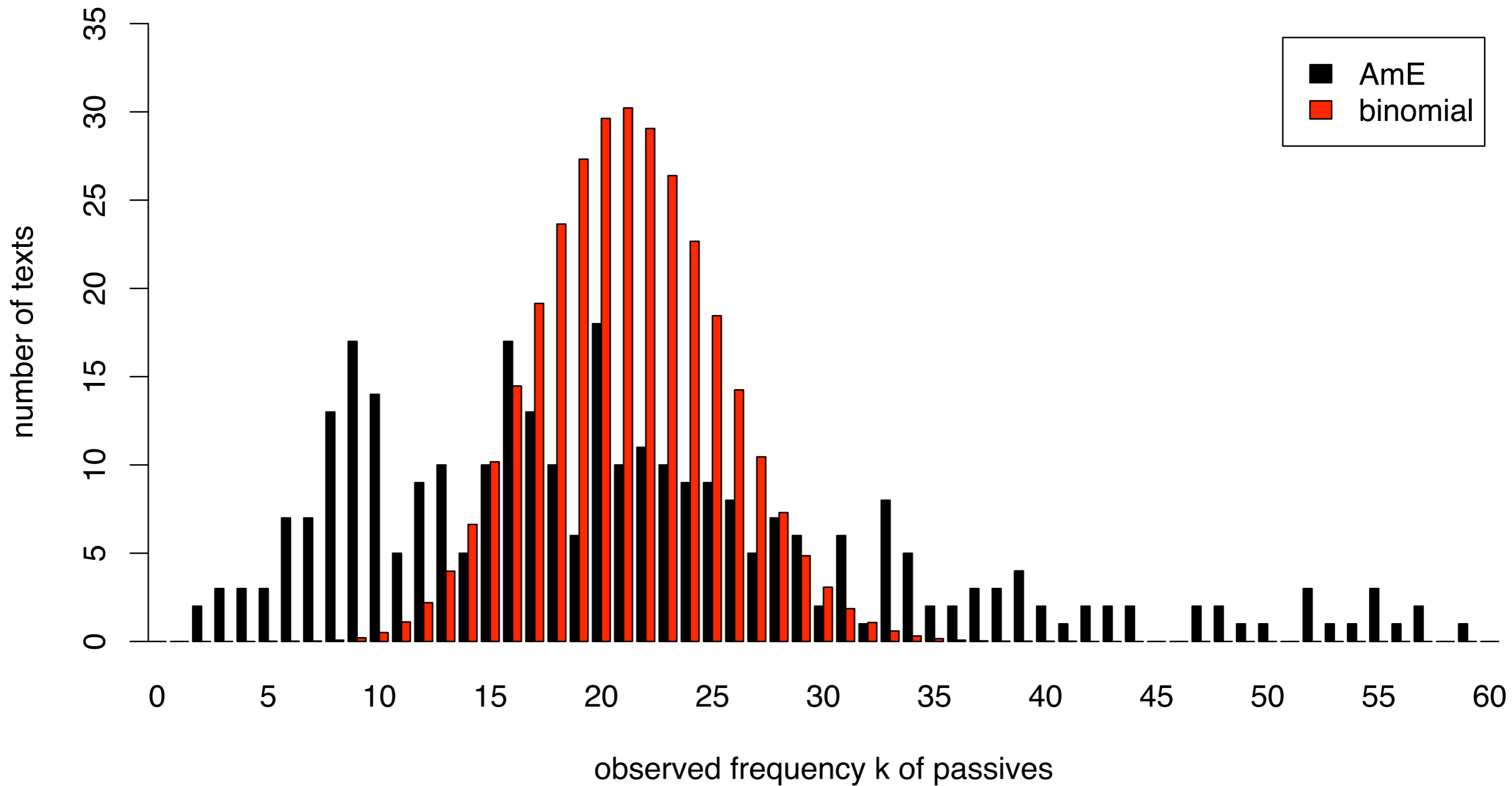
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- ◆ No we know how to measure non-randomness
 - find out if corpus texts are random samples
 - i.e., if they follow a binomial sampling distribution
 - ➔ tabulate observed frequencies across corpus texts

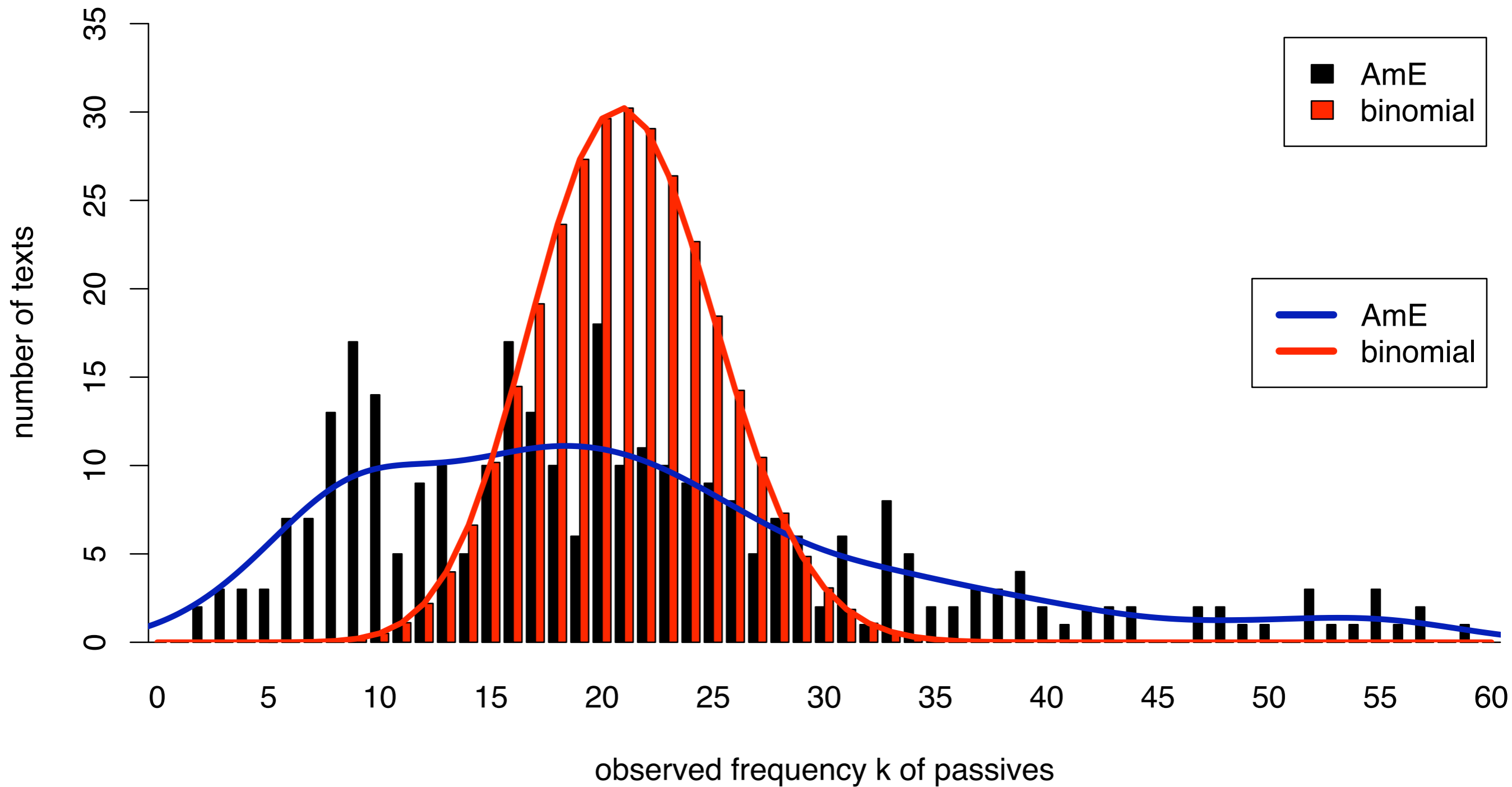
Measuring non-randomness

- ◆ Tabulate number of texts with k passives
 - illustrated for subsets of Brown/LOB (310 texts each)
 - meaningful because all texts have the same length
- ◆ Compare with binomial distribution
 - for population proportion $H_0 : \pi = 21.1\%$ (Brown) and $\pi = 22.2\%$ (LOB); approx. $n = 100$ sentences per text
 - estimated from full corpus \rightarrow best possible fit
- ◆ Non-randomness \rightarrow larger sampling variation

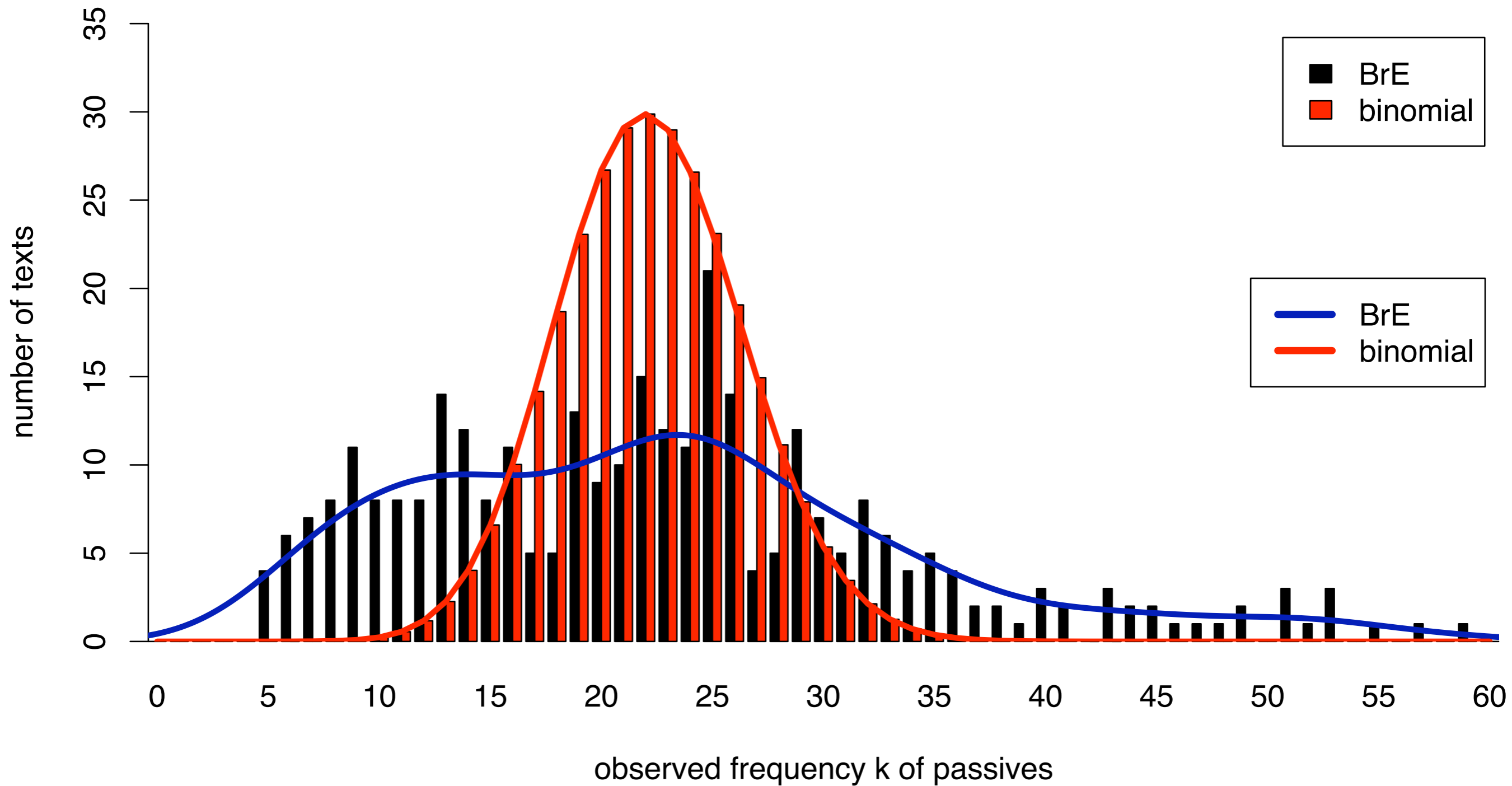
Passives in the Brown corpus

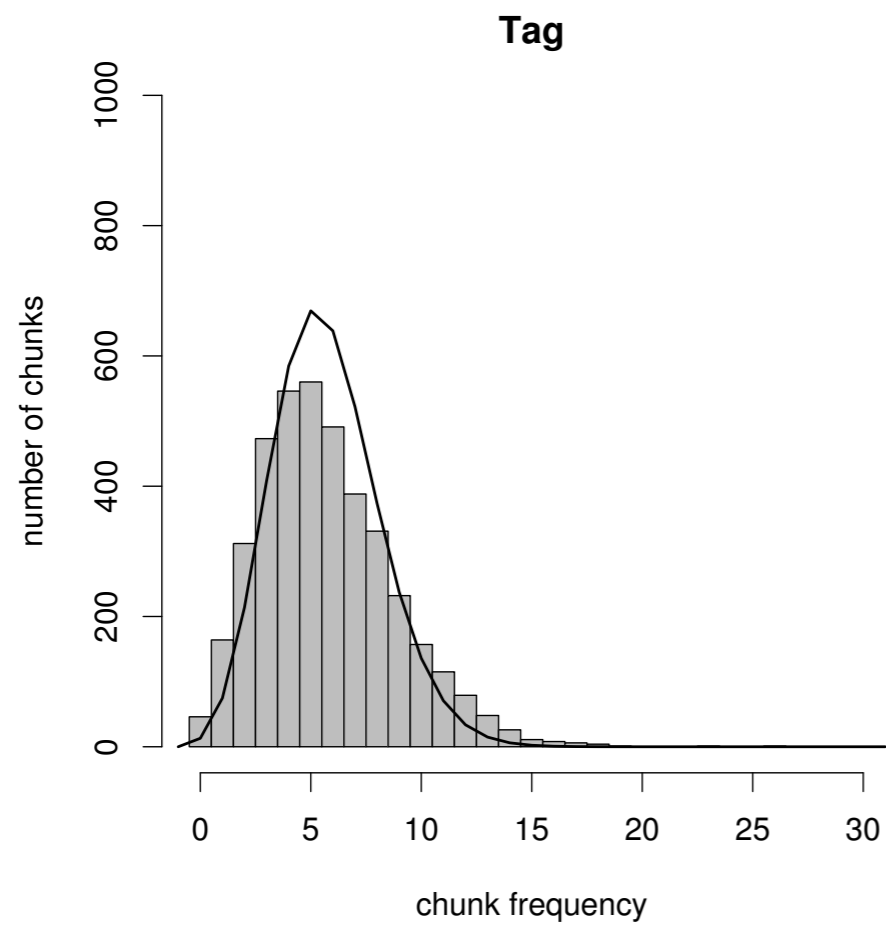


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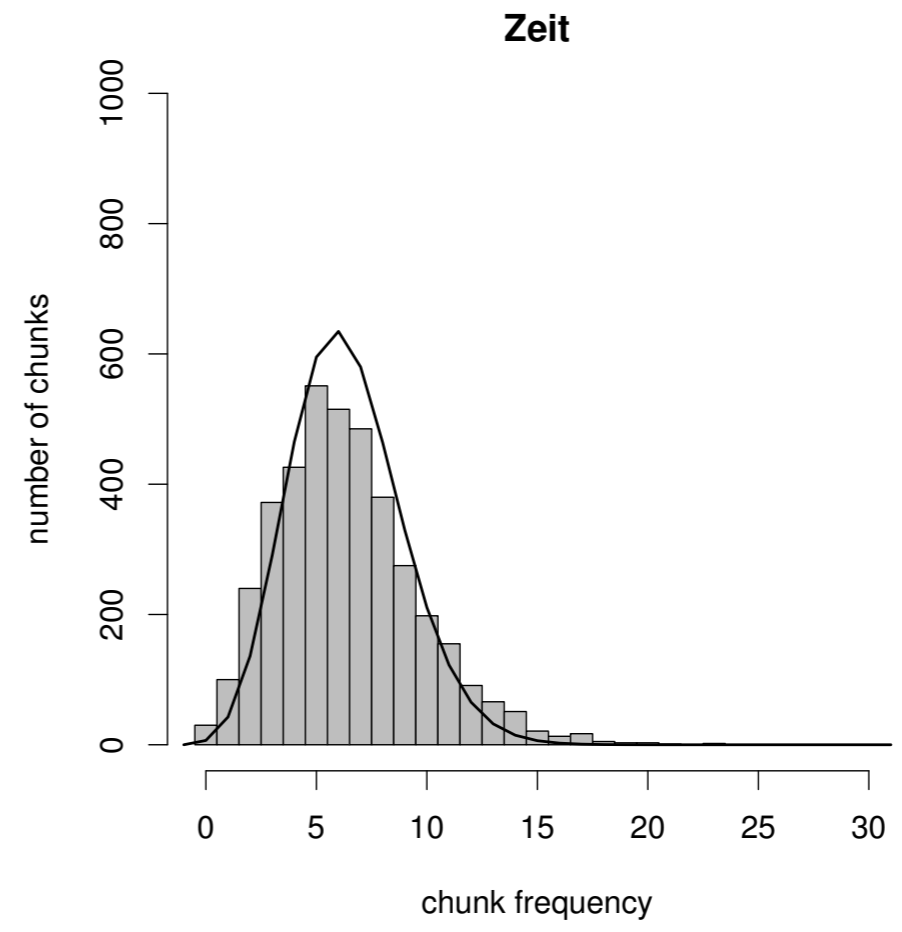
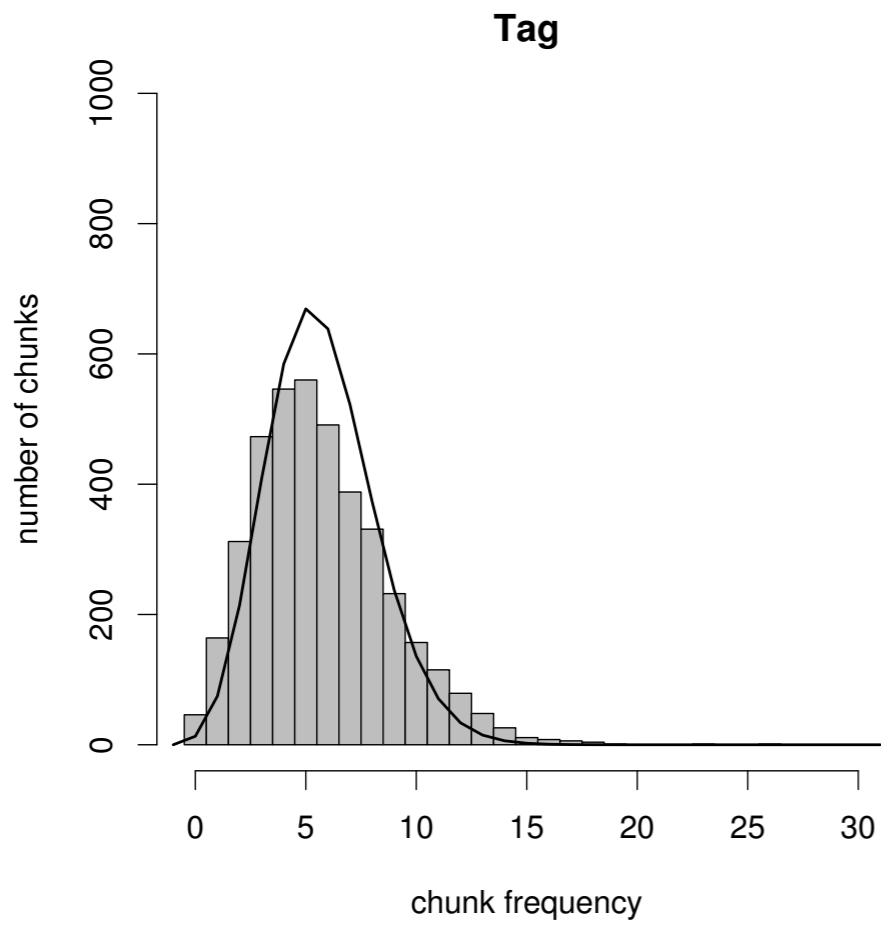


Passives in the LOB corpus

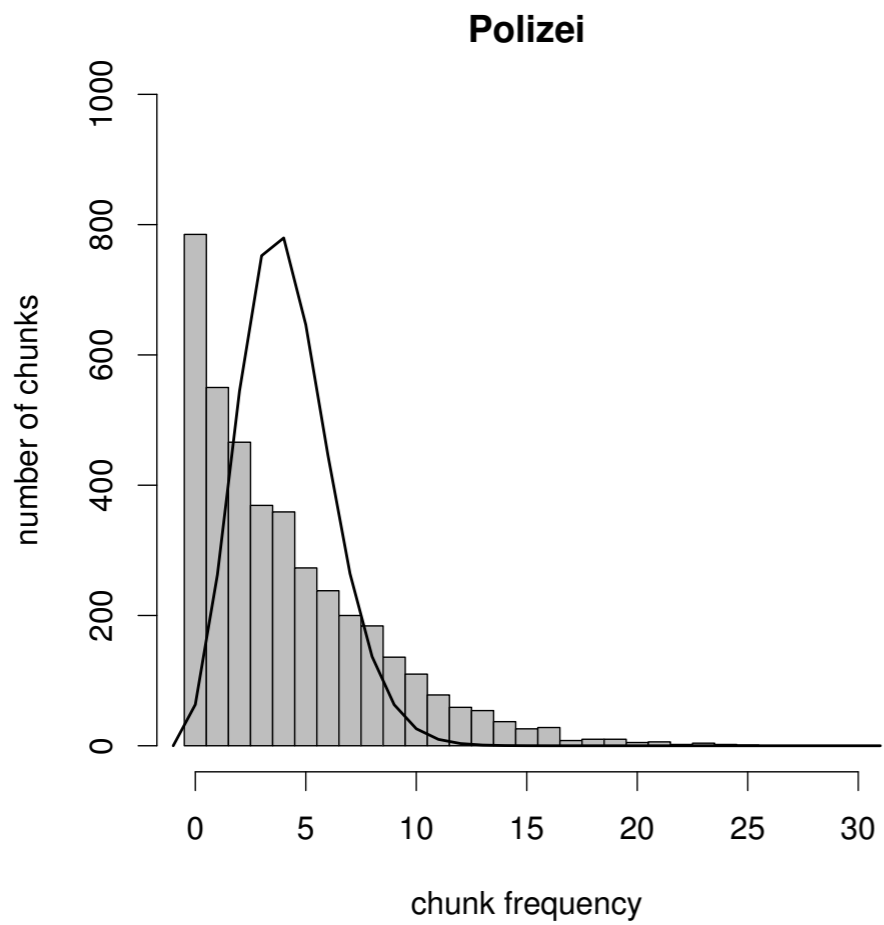
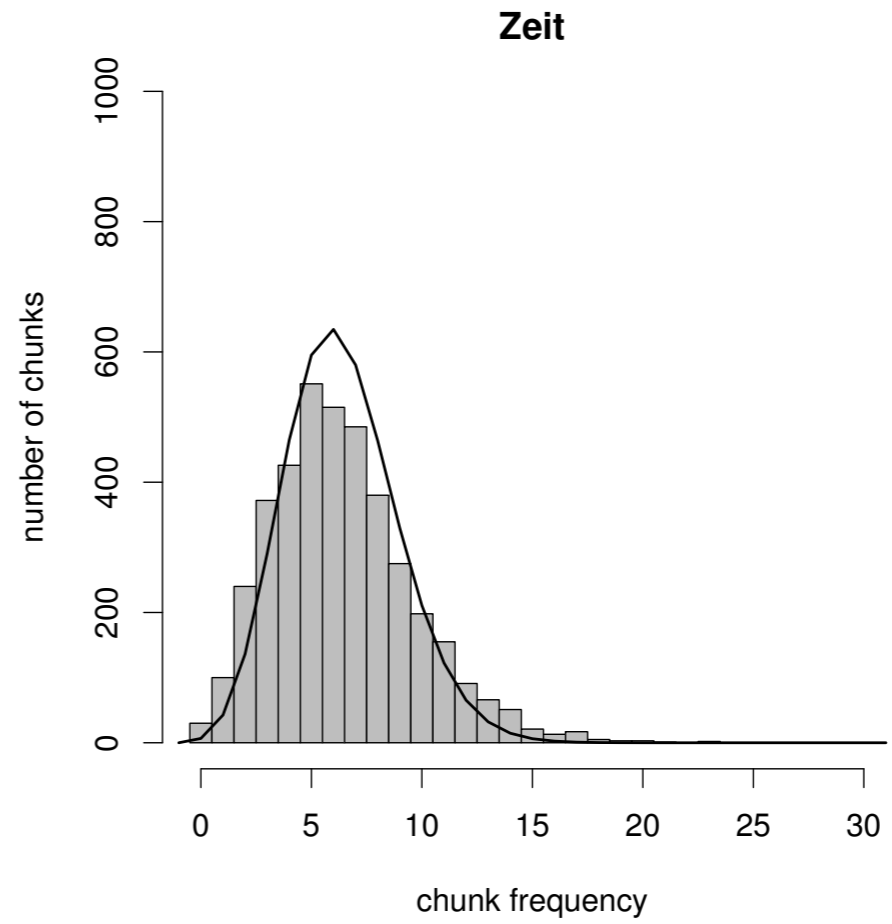
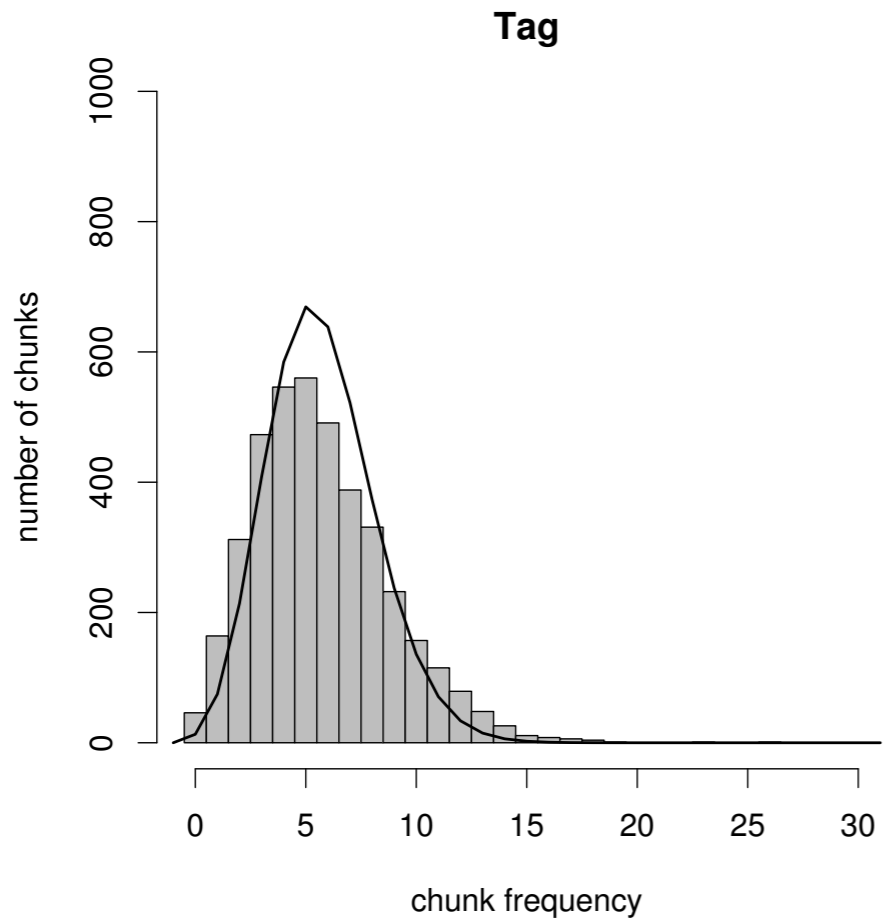




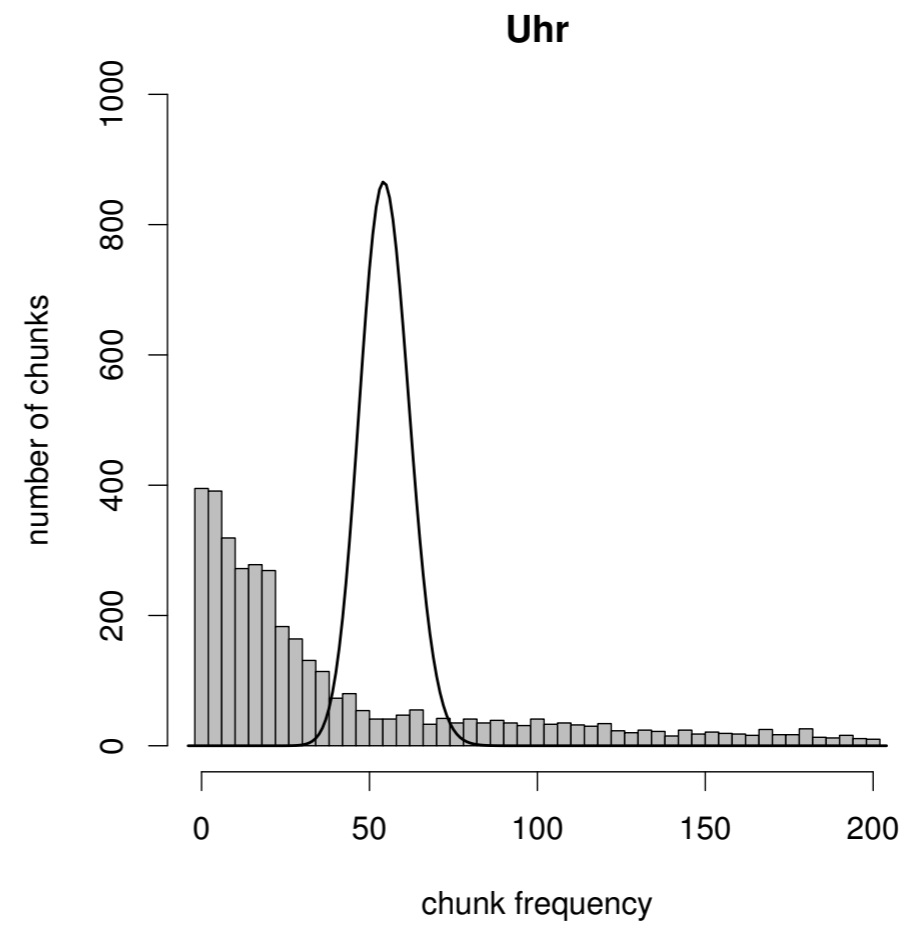
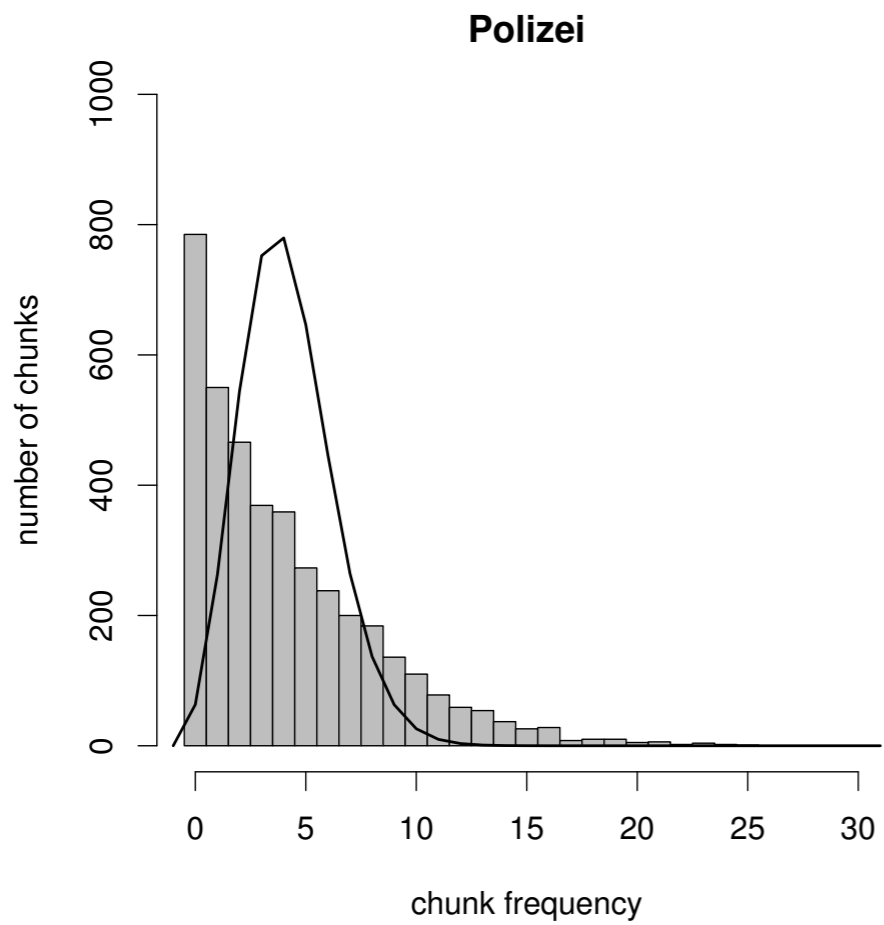
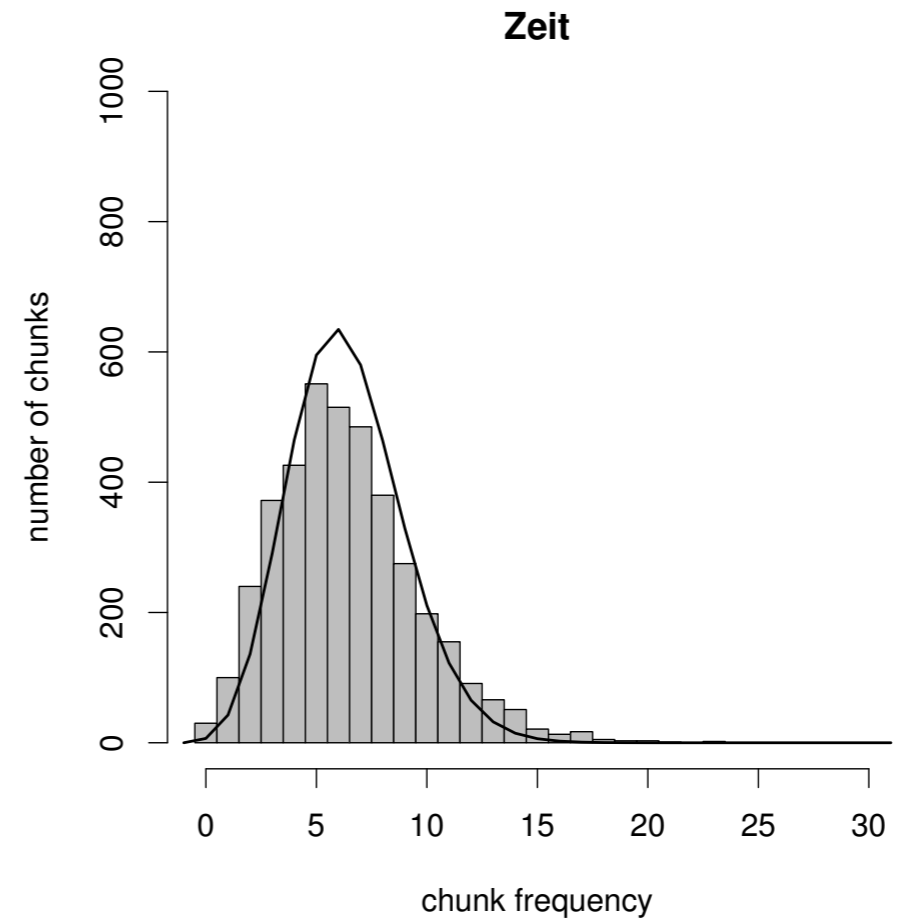
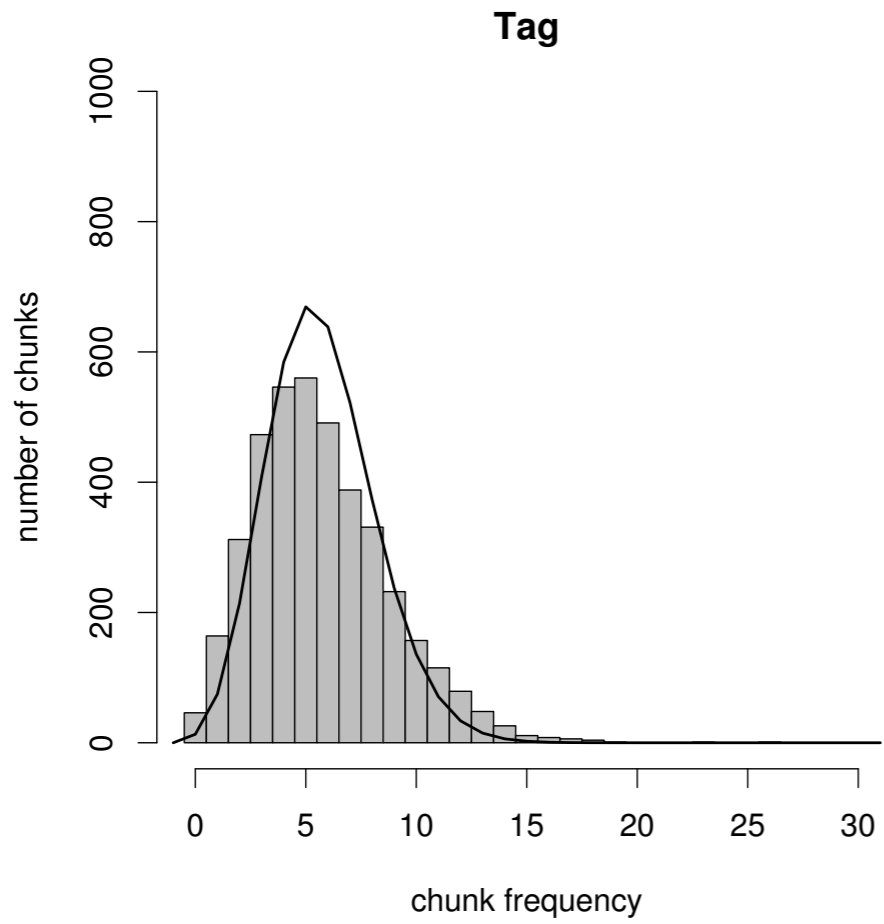
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 - results in much smaller sample size ...
(BNC: 4,048 texts rather than 6,023,627 sentences)
 - ... but more informative measurements (relative frequencies on interval rather than nominal scale)

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(BNC: 4,048 texts rather than 6,023,627 sentences)
 - ... but more informative measurements (relative frequencies on interval rather than nominal scale)
- ◆ Use statistical techniques that account for the **overdispersion** of relative frequencies
 - Gaussian distribution allows us to estimate **spread** (variance) independently from **location**
 - Standard technique: **Student's t-test**

A case study: Passives in AmE and BrE

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- ◆ Are there more passives in BrE than in AmE?
 - based on data from subsets of Brown and LOB
 - 9 categories: press reports, editorials, skills & hobbies, misc., learned, fiction, science fiction, adventure, romance
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 - AmE: 6584 out of 31,173 sentences = 21.1%
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- ◆ Chi-squared test (→ pooled data, binomial)
vs. t-test (→ sample of texts, Gaussian)

Let's do that in R ...

```
# passive counts for each text in Brown and LOB corpus
> Passives <- read.delim("passives_by_text.tbl")

# display 10 random rows to get an idea of the table layout
> Passives[sample(nrow(Passives), 10), ]

# add relative frequency of passives in each file (as percentage)
> Passives <- transform(Passives,
  relfreq = 100 * passive / n_s)

# split into separate data frames for Brown and LOB texts
> Brown <- subset(Passives, lang=="AmE")
> LOB <- subset(Passives, lang=="BrE")
```

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- ◆ Chi-squared test: **highly significant**
 - p-value: **.00069** < .001
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- ◆ R code: pooled counts + proportions test

```
> passives.B <- sum(Brown$passive)
> n_s.B <- sum(Brown$n_s)
> passives.L <- sum(LOB$passive)
> n_s.L <- sum(LOB$n_s)

> prop.test(c(passives.L, passives.B),
            c(n_s.L, n_s.B))
```


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- ◆ t-test: **not significant**
 - p-value: **.1340** > .05 ($t=1.50$, $df=619.96$)
 - confidence interval for difference: **-0.6% – +4.9%**
 - H_0 : same average relative frequency in AmE and BrE

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 - H_0 : same average relative frequency in AmE and BrE
- ◆ R code: apply `t.test()` function
 - > `t.test(LOB$relfreq, Brown$relfreq)`
 - # alternative syntax: “formula” interface**
 - > `t.test(relfreq ~ lang, data=Passives)`



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 - 50% written / 50% spoken: $\pi = 13.0\%$

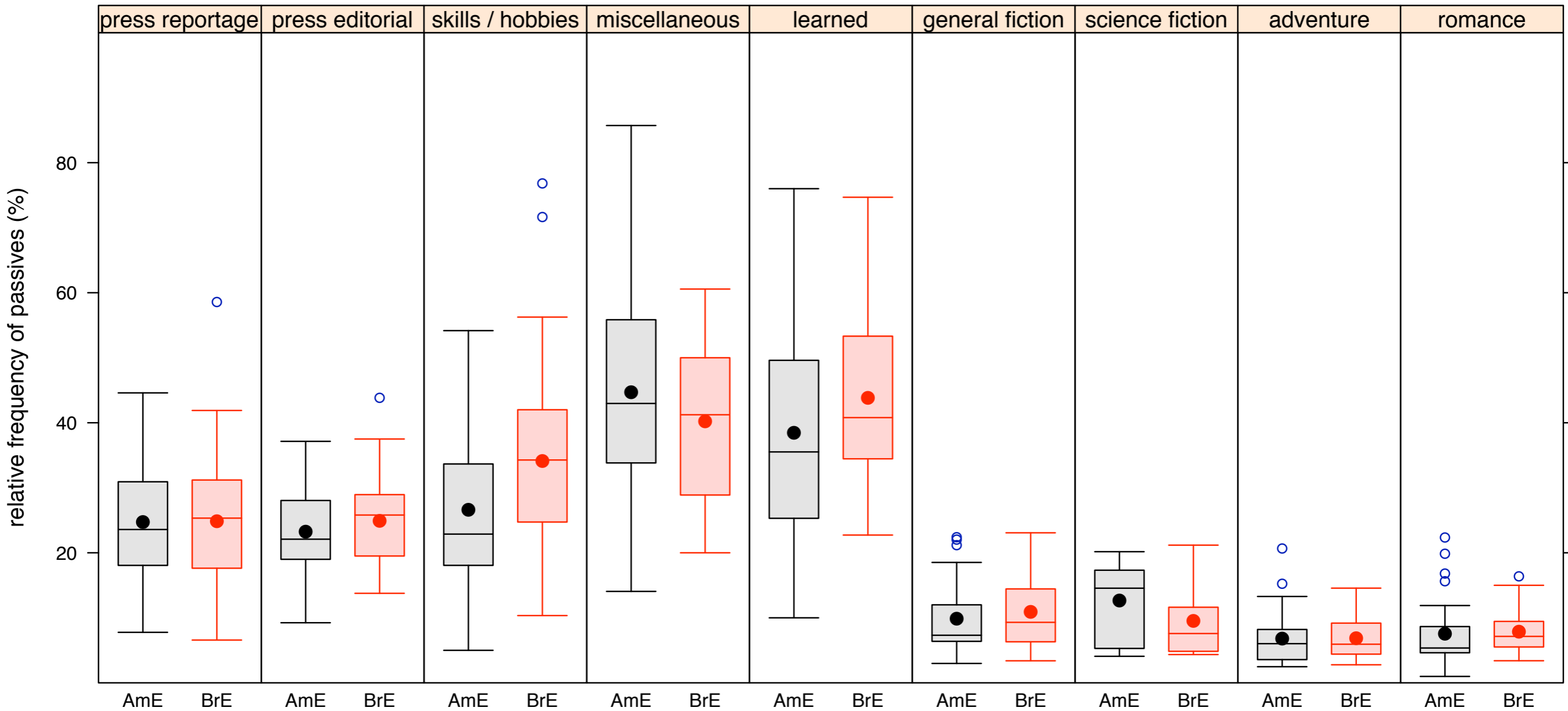
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 - 90% written / 10% spoken: $\pi = 16.6\%$

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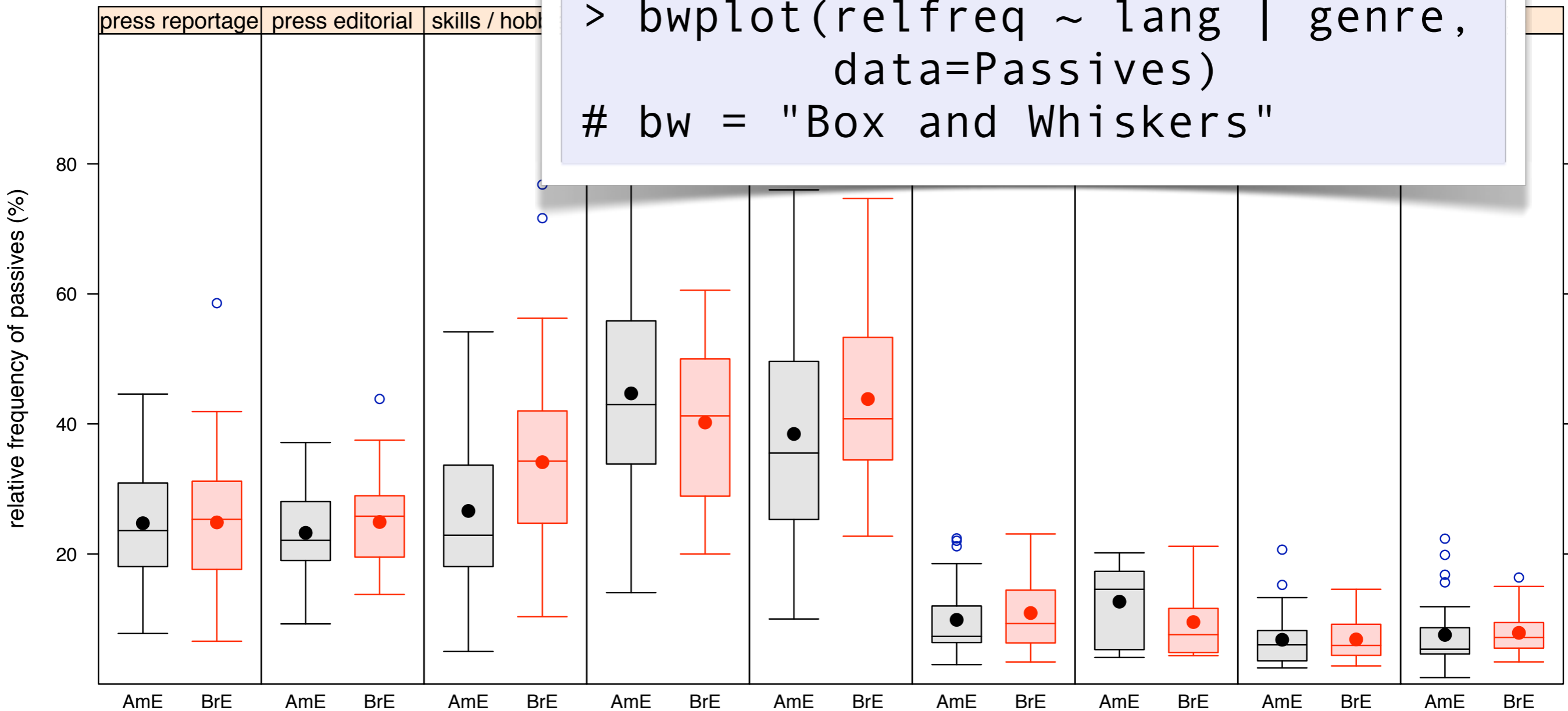
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 - 20% written / 80% spoken: $\pi = 6.8\%$

Average relative frequency?

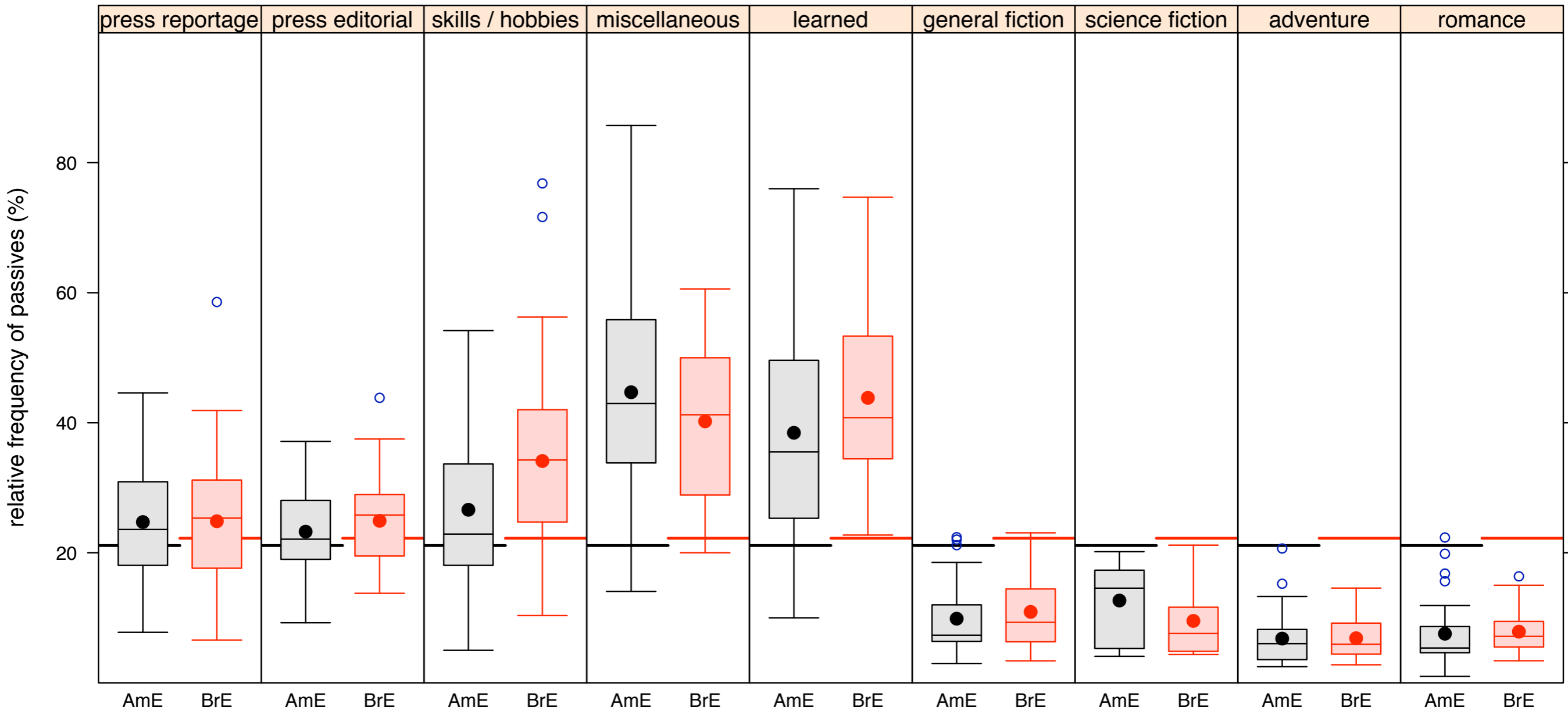


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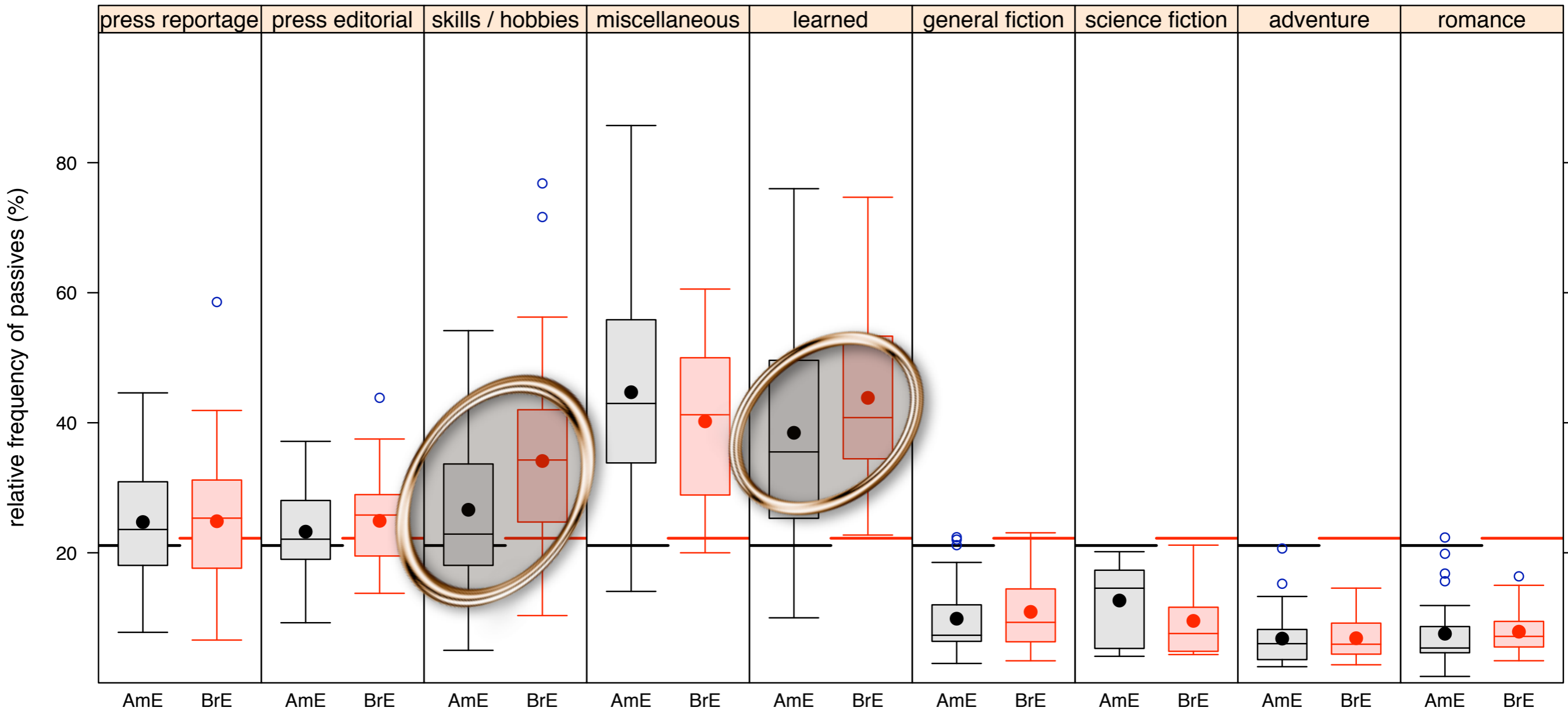
```
> library(lattice)
> bwplot(relfreq ~ lang | genre,
         data=Passives)
# bw = "Box and Whiskers"
```



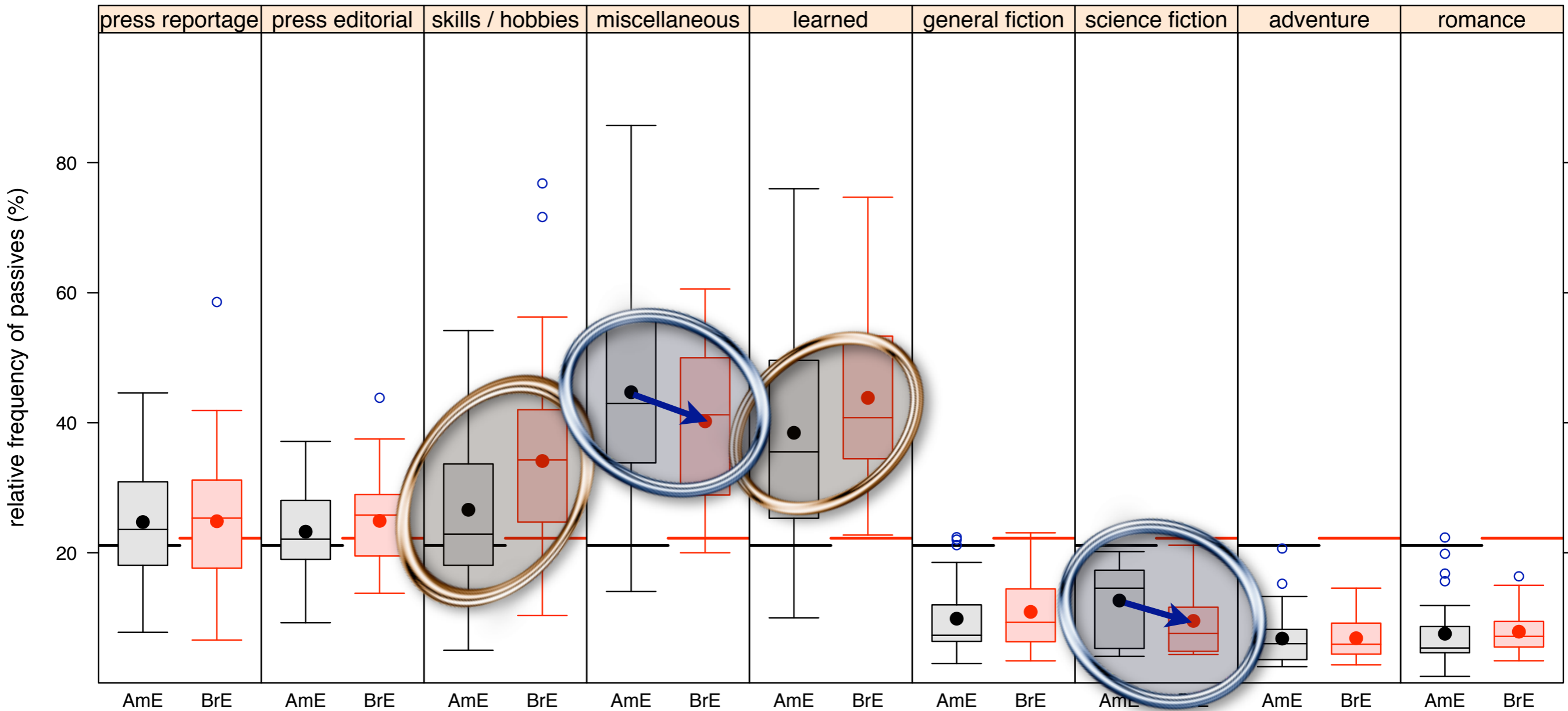
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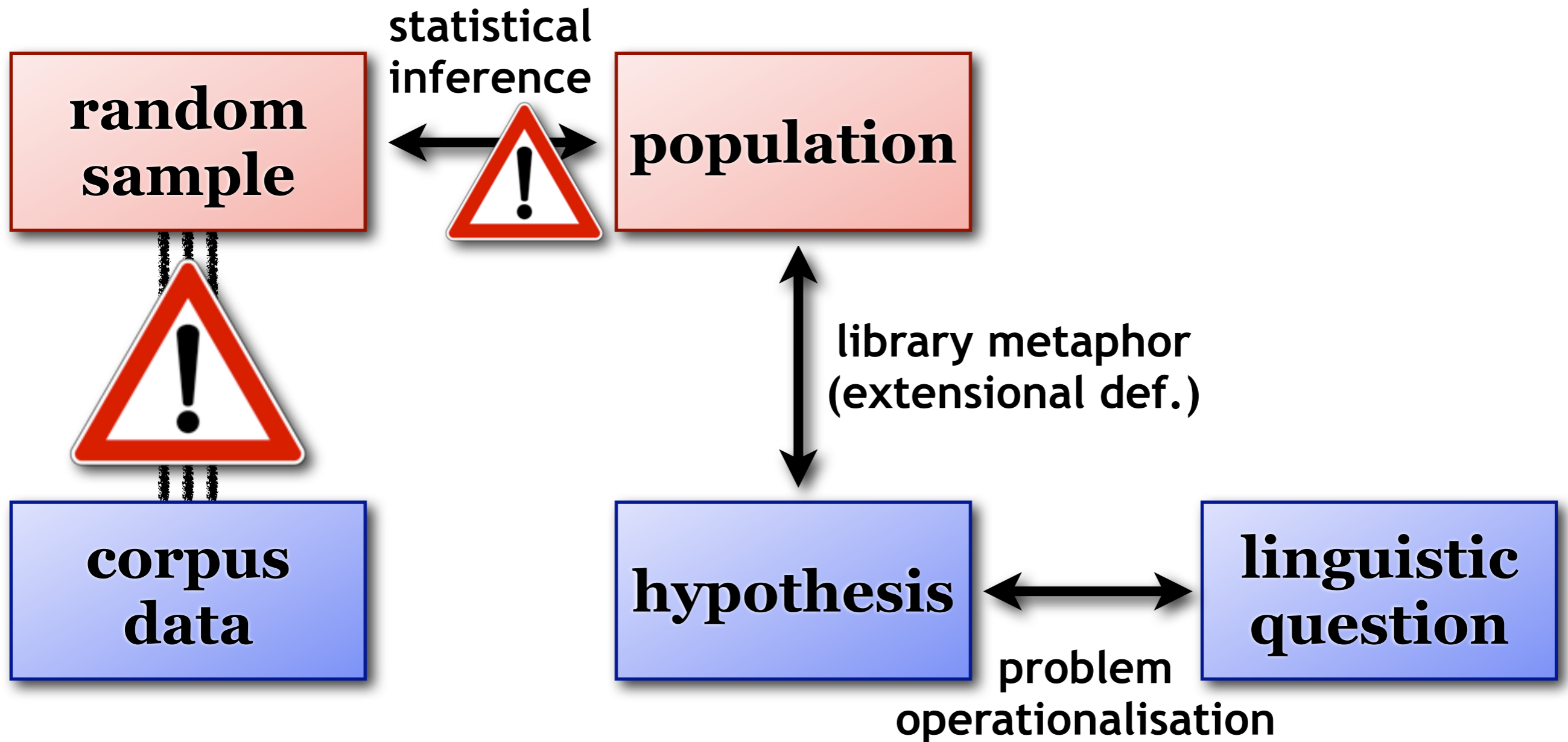
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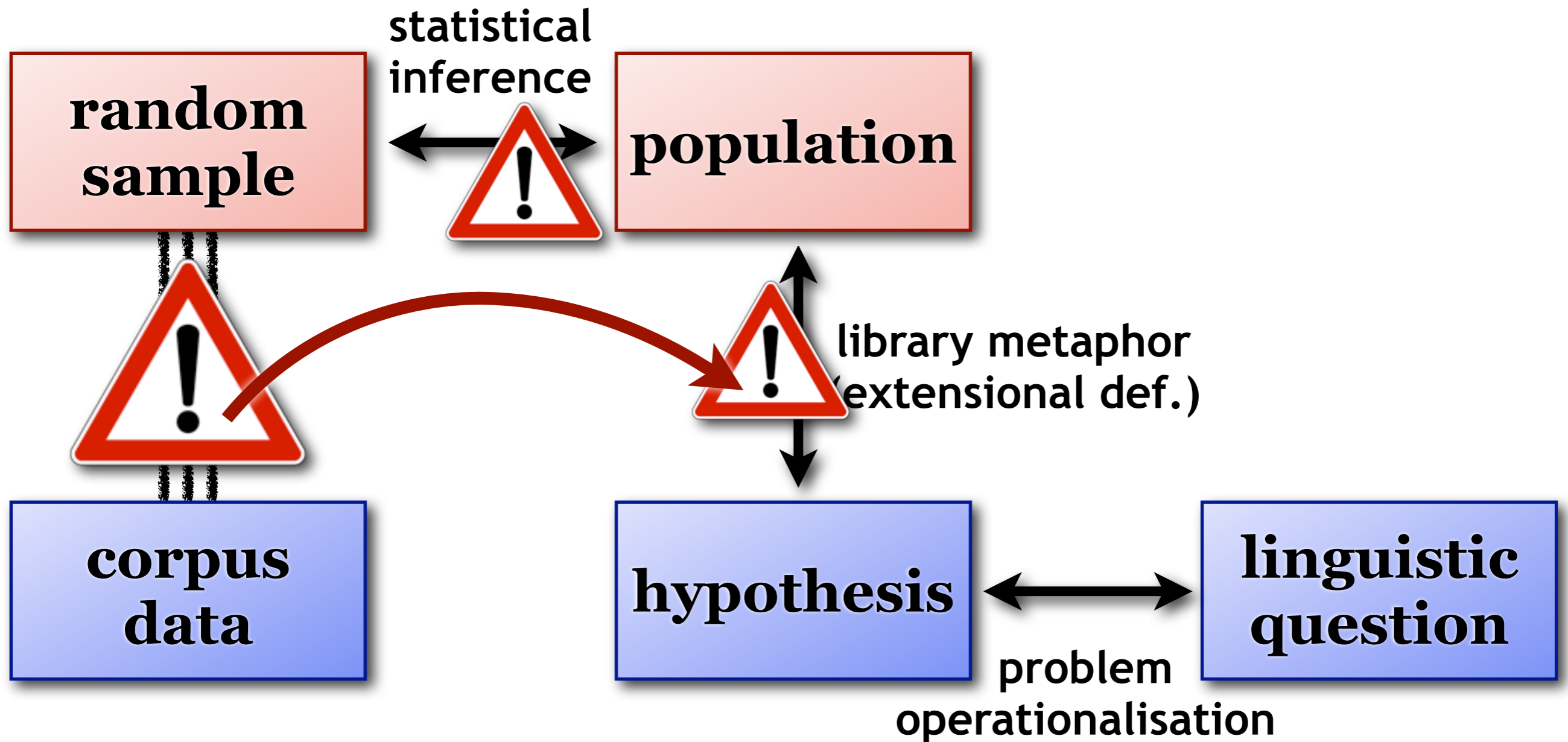
Average relative frequency?



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Rethinking corpus frequencies

Studying variation in language

- ◆ It seems absurd now to measure & compare relative frequencies in “language” (= library)
 - proportion π depends more on composition of library than on properties of the language itself
- ◆ Quantitative corpus analysis has to account for the variation of relative frequencies between individual texts (cf. Gries 2006)
 - research question → one factor behind this variation

Studying variation in language

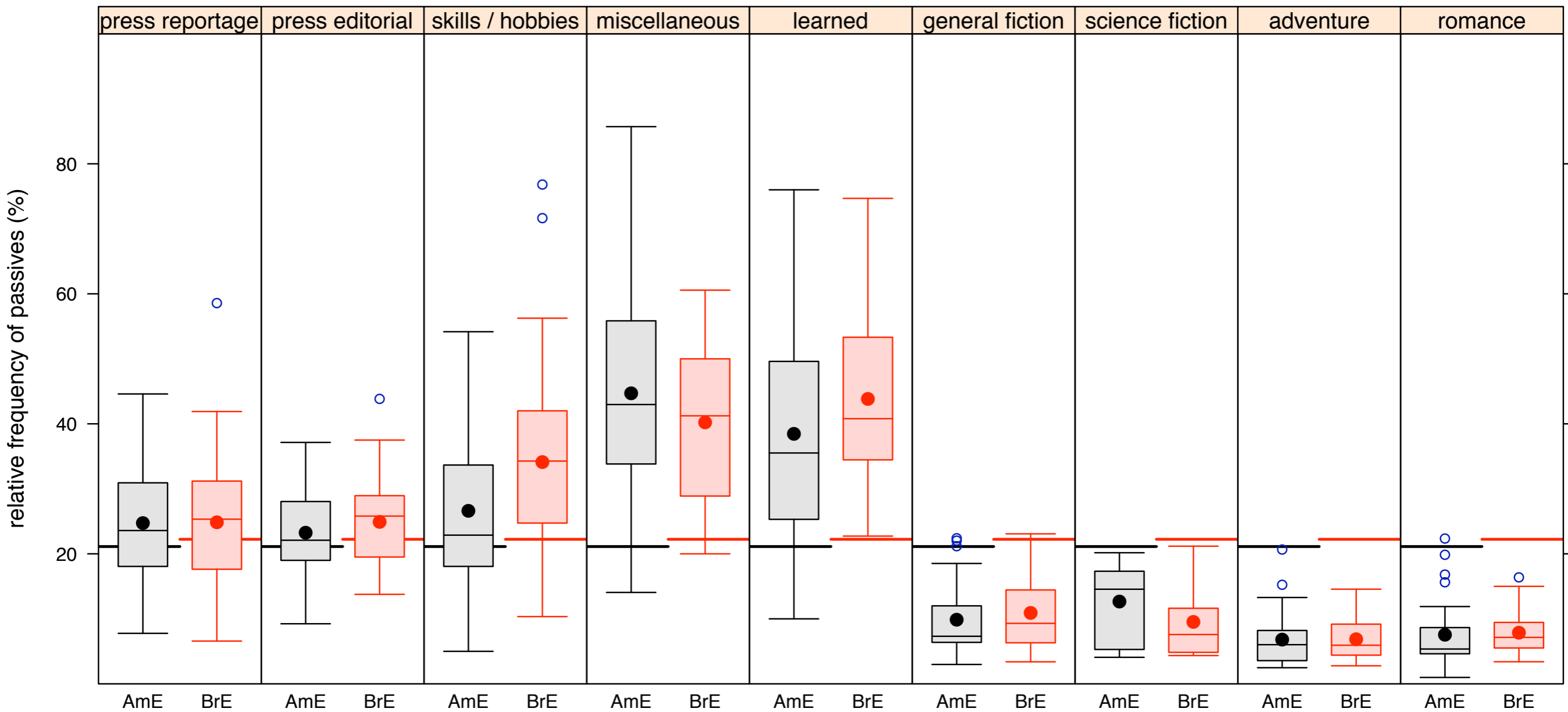
Studying variation in language

- ◆ **Approach 1:** restrict study to sublanguage in order to eliminate non-randomness
 - data from this sublanguage (= single section in library) can be pooled into large random sample

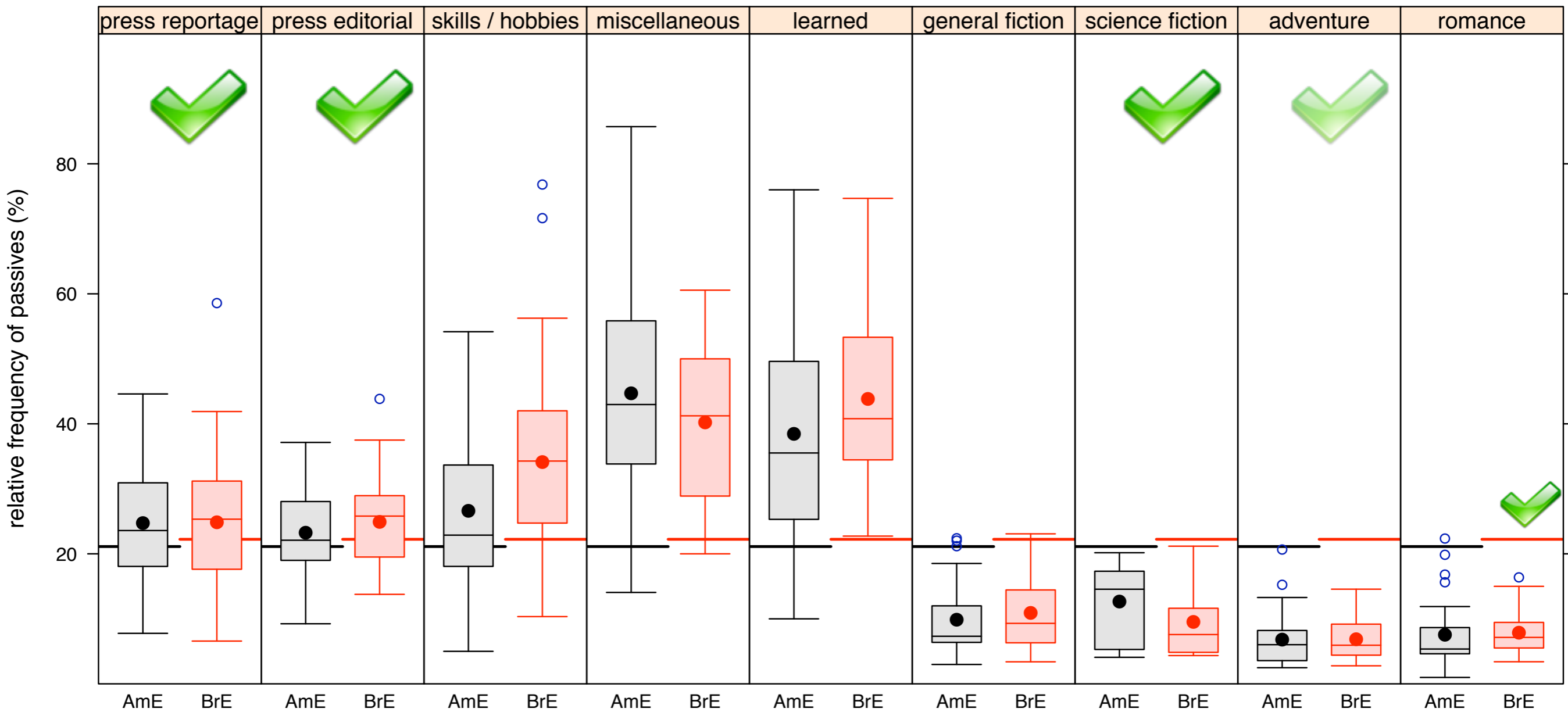
Studying variation in language

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 - data from this sublanguage (= single section in library) can be pooled into large random sample
- ◆ **Approach 2:** goal of quantitative corpus analysis is to **explain variation** between texts in terms of
 - random sampling (of tokens within text)
 - stylistic variation: genre, author, domain, register, ...
 - subject matter of text → term clustering effects
 - differences between language varieties ← **research question**

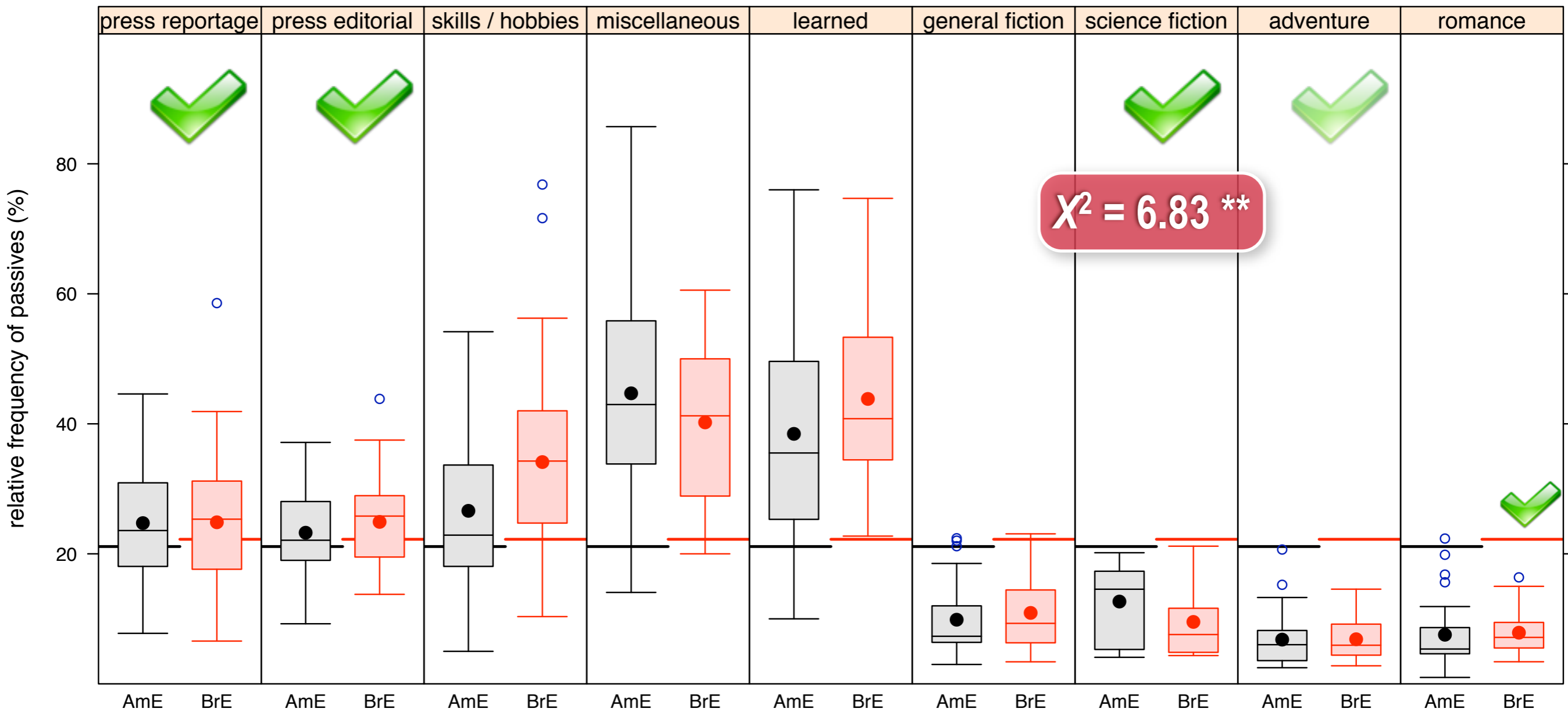
Eliminating non-randomness



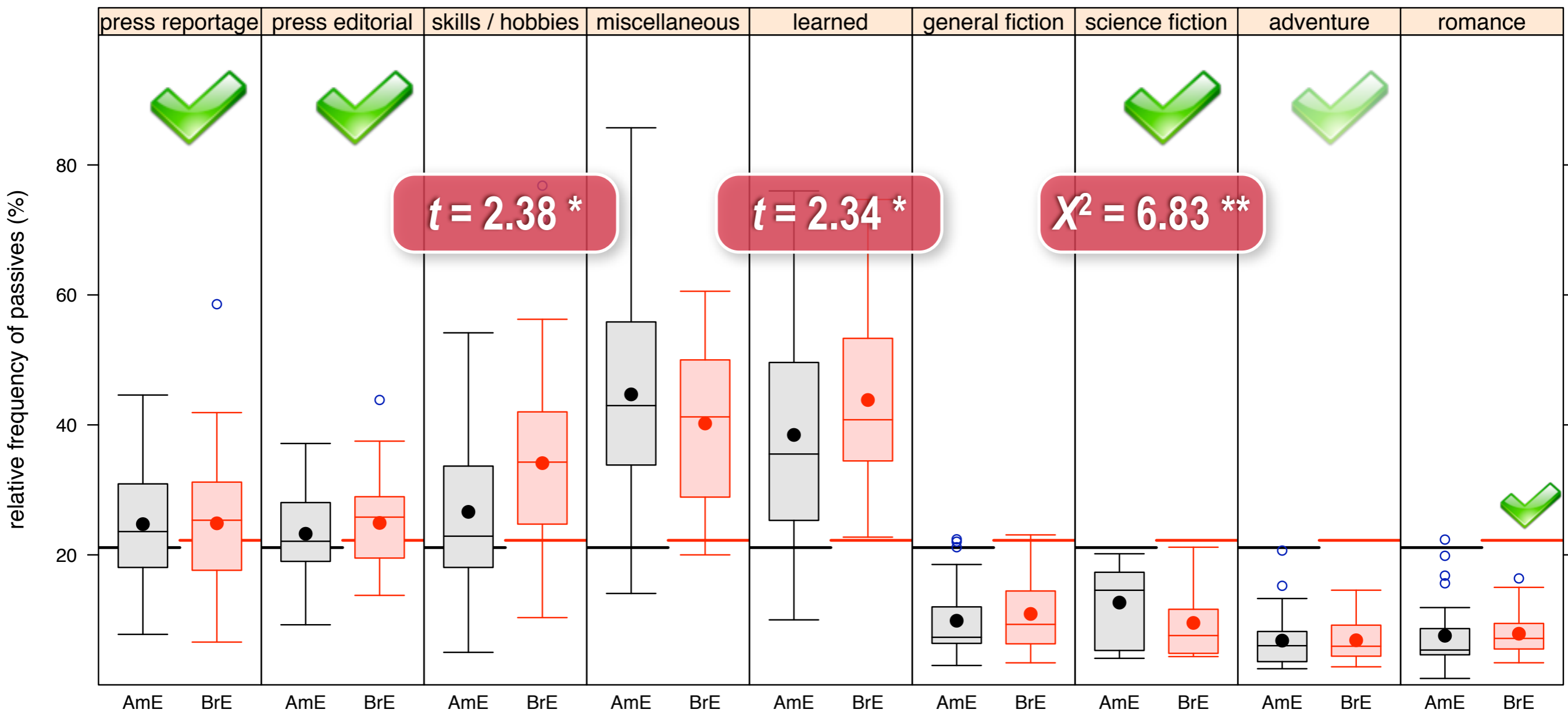
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Explaining variation

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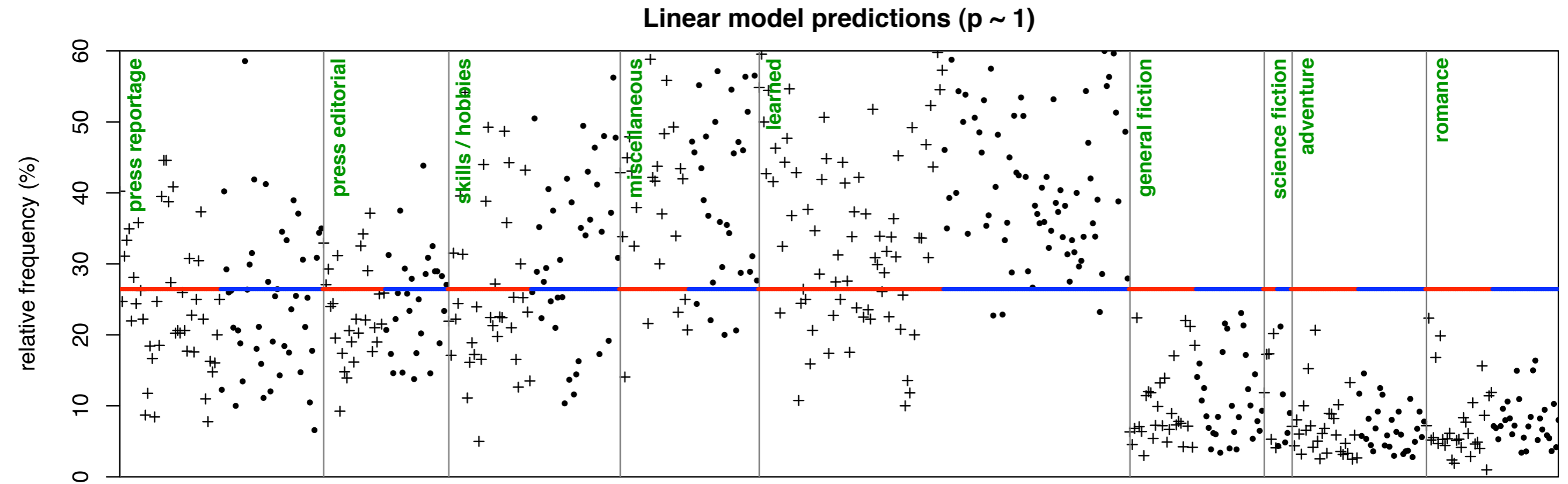
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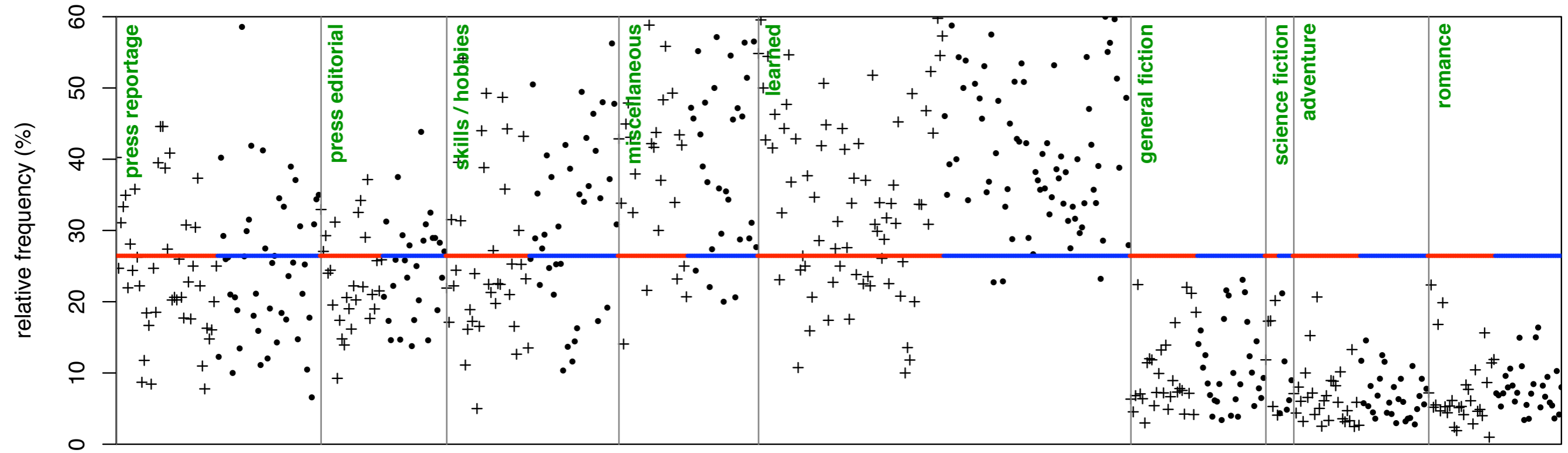
I’m just an ANOVA ...

Linear model for passives

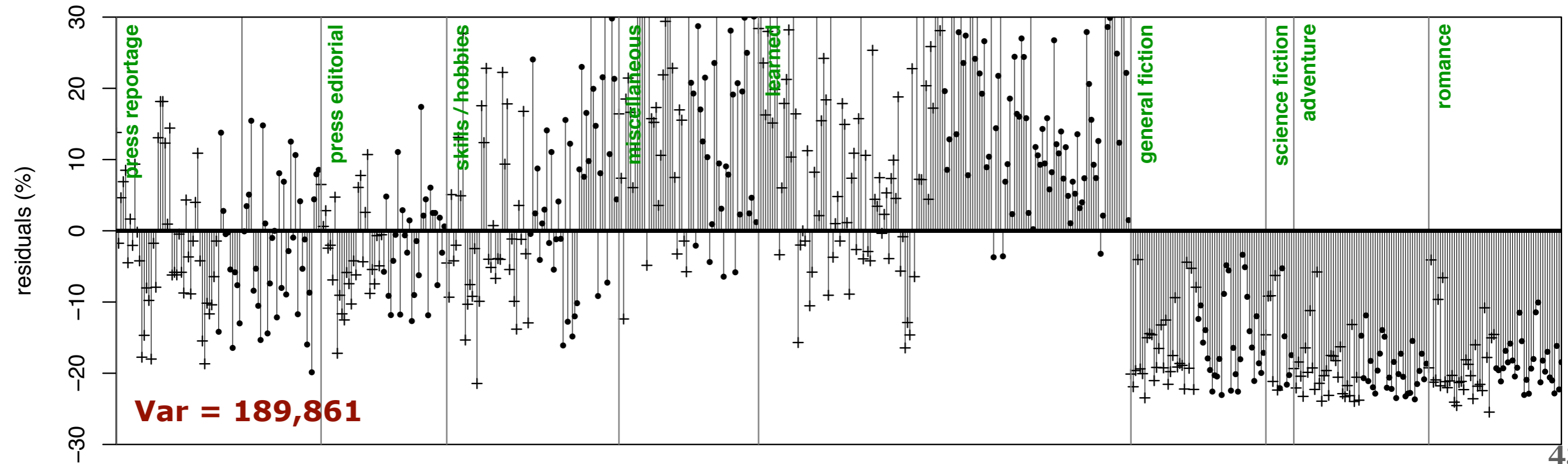


Linear model for passives

Linear model predictions ($p \sim 1$)

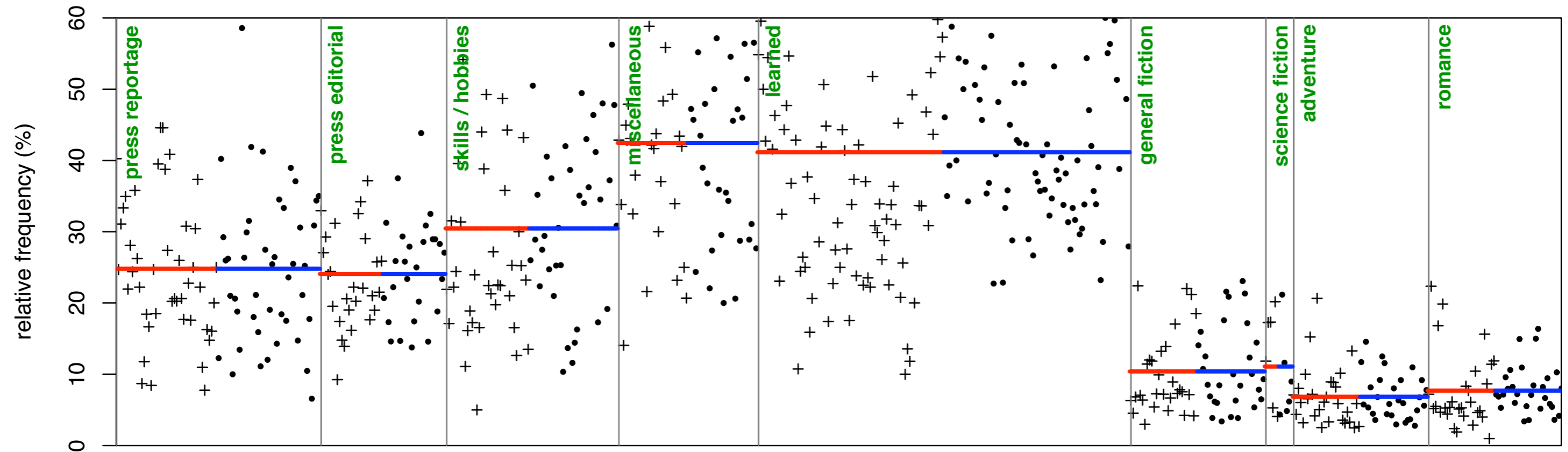


Unexplained residuals of linear model

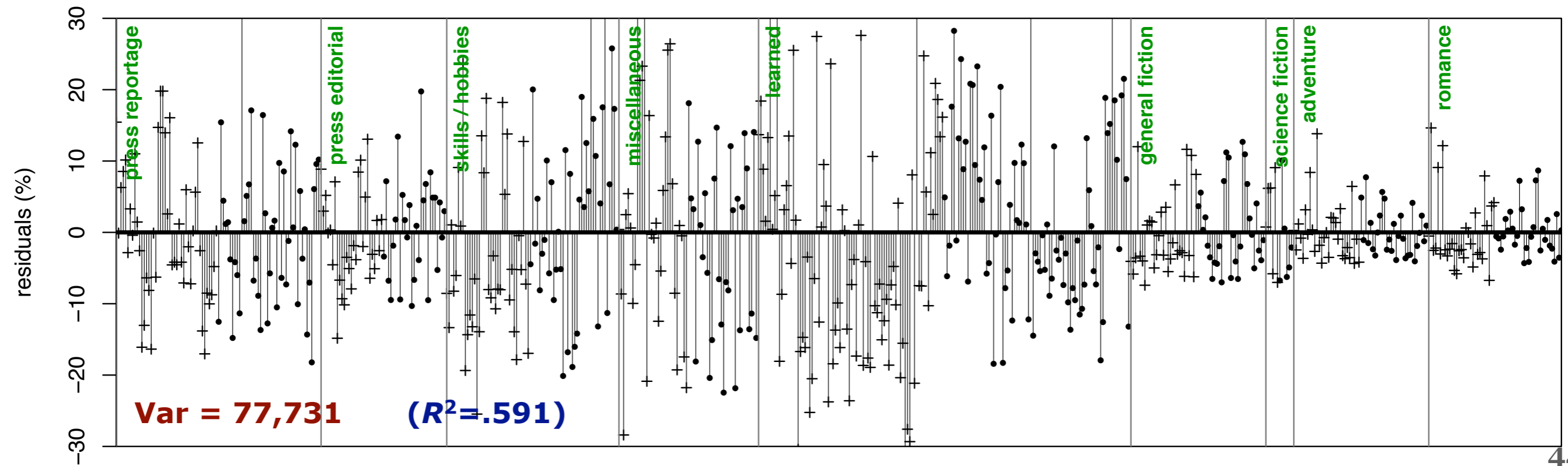


Linear model for passives

Linear model predictions ($p \sim 1 + \text{genre}$)

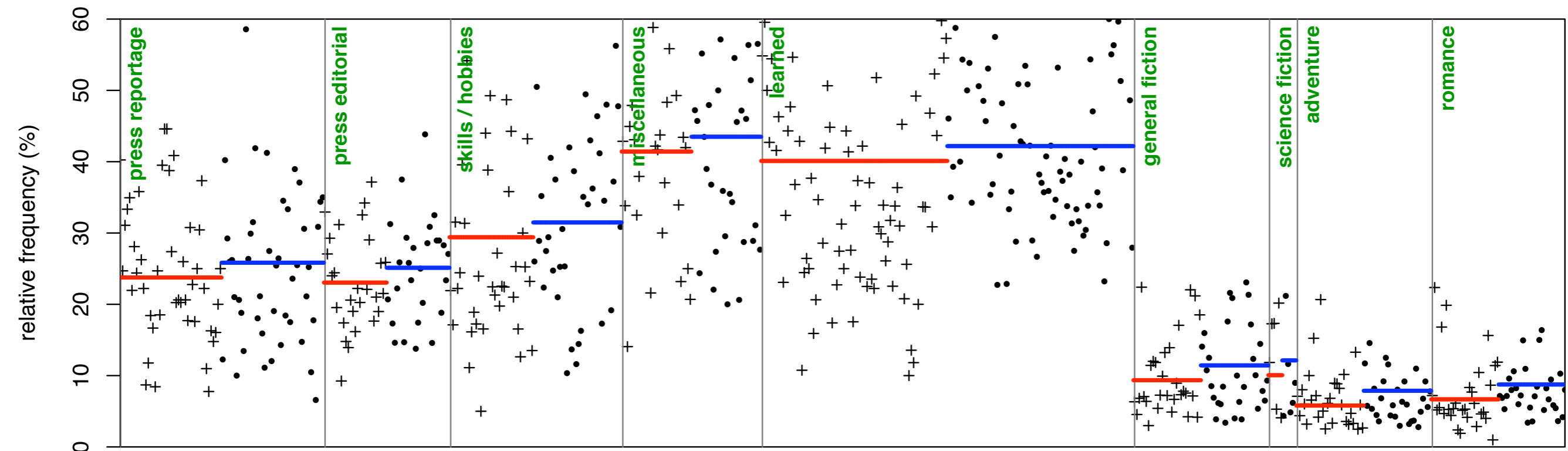


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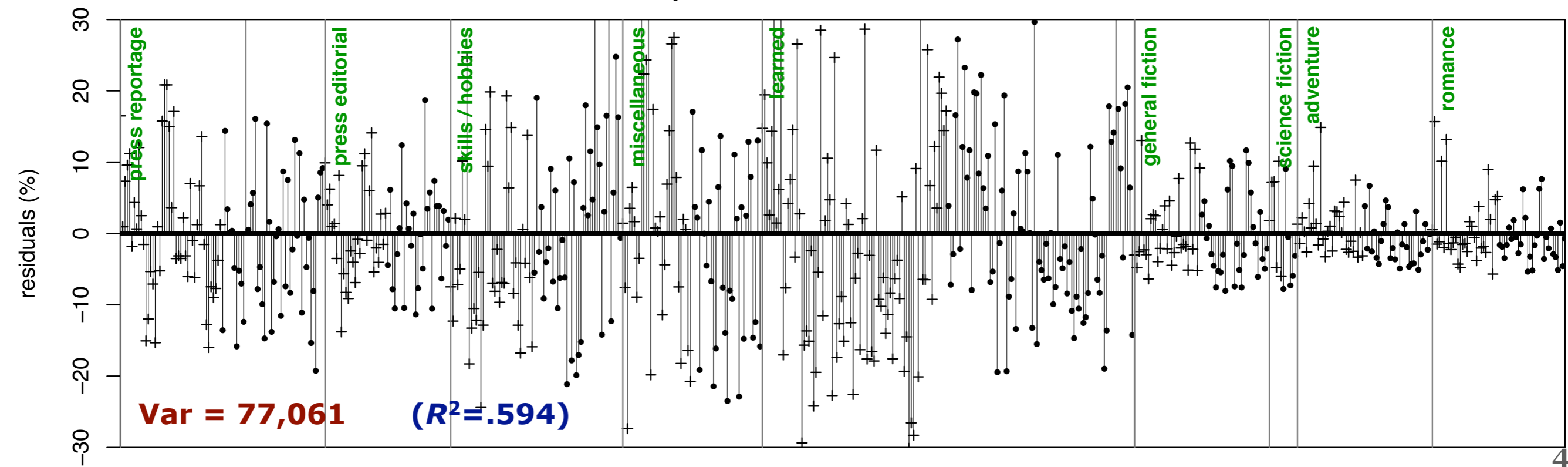


Linear model for passives

Linear model predictions ($p \sim 1 + \text{genre} + \text{Am/Br}$)



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- binomial sampling variation: ca. 10,200 (= 5.4%)

Linear models in R

```
# linear model "formula": response ~ explanatory factors
# (here, only main effects without genre/language interaction)
> LM <- lm(relfreq ~ genre + lang, data=Passives)

# analysis of variance shows which factors are significant
> anova(LM)      # see ?anova.lm for details

# individual coefficients + standard errors
> summary(LM)
> confint(LM)    # corresponding confidence intervals

# interaction term improves model fit, but is not quite significant
> LM <- lm(relfreq ~ genre + lang + genre:lang,
           data=Passives)
> anova(LM)
```

Linear model for passives

- ◆ F-tests show significant effects of genre ($p < 10^{-15}$) and AmE / BrE ($p = .0198$)
- ◆ 95% confidence intervals for effect sizes:
 - AmE / BrE: 0.3% ... 3.8%
 - genre = learned 13.4% ... 19.3%
 - compared to “press reportage” genre as baseline
 - genre = romance -20.8% ... -13.4%
 - genre = ...

Linear models in R

```
# more intuitive than coefficients: model predictions for each
# genre and language variety; based on “dummy” data frame with
# all possible genre/language combinations (ordered by genre)
> Predictions <- unique(
  Passives[, c("genre", "lang")])
> Predictions <- Predictions[
  order(Predictions$genre, Predictions$lang), ]

# predicted average relative frequency of passives in each category
> transform(Predictions,
  predicted=predict(LM, newdata=Predictions))

# confidence and prediction intervals
> cbind(Predictions, predict(LM,
  newdata=Predictions, interval="confidence"))
> cbind(Predictions, predict(LM,
  newdata=Predictions, interval="prediction"))
```



Linear models are not appropriate!

```
> par(mfrow=c(2,2))  
> plot(LM)  
> par(mfrow=c(1,1))
```



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- ◆ Predictions not restricted to range 0% – 100%

Generalised linear models

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$$f_i \sim B(n_i, \pi_i)$$

binomial sampling
("family")

$$\pi_i = \frac{1}{1 + e^{-\theta_i}}$$

"link" function

linear predictor $\rightarrow \theta_i = \beta_0 + \beta_1(\text{genre}) + \beta_2(\text{AmE/BrE})$

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◆ Interpretation of confidence intervals difficult

GLM in R

(note the extra options needed!)

```
# for GLM with binomial family, responses are pairs of
# passive / active counts ( $f_k, n_k - f_k$ ) = "successes" / "failures"
> response.matrix <- cbind(Passives$passive,
  Passives$n_s - Passives$passive)

# genre * lang is shorthand for main effects + all interactions
> GLM <- glm(response.matrix ~ genre * lang,
  family="binomial", data=Passives)

# individual coefficients + standard errors
> anova(GLM, test="Chisq") # interaction significant now
> summary(GLM) # even more difficult to interpret than for LM
> confint(GLM)

# diagnostics plot (; separate multiple commands in single line)
> par(mfrow=c(2,2)); plot(GLM); par(mfrow=c(1,1))
```

GLM in R

(note the extra options needed!)

predictions for each genre and language variety

```
> transform(Predictions, predicted = 100 *  
  predict(GLM, type="response",  
  newdata=Predictions))
```

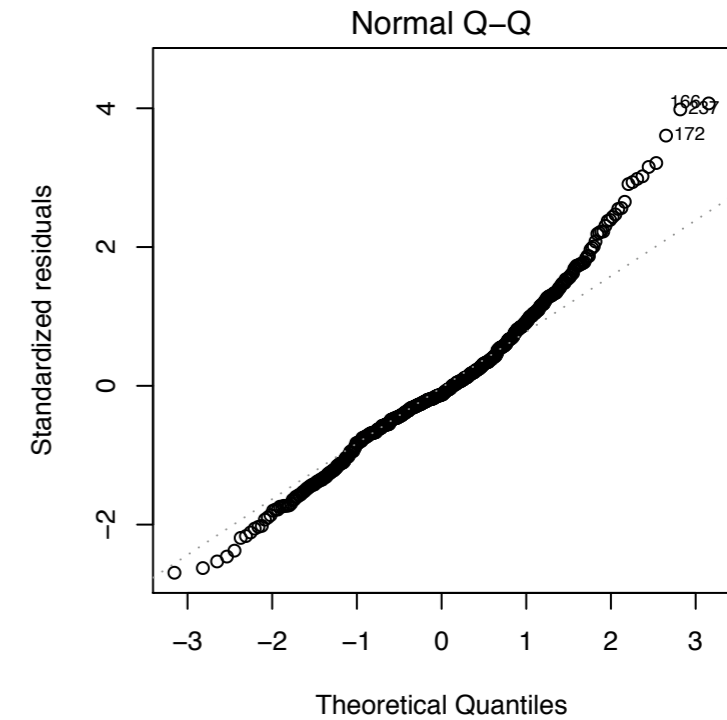
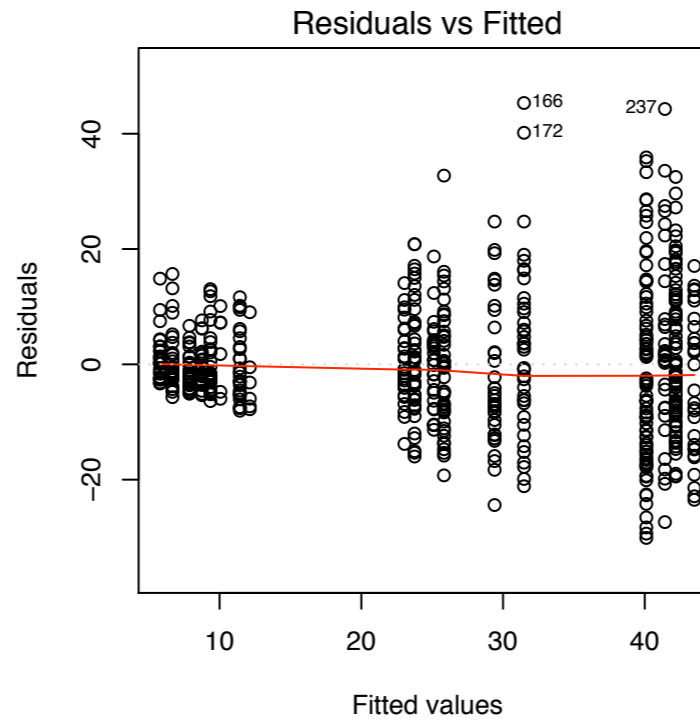
calculate confidence intervals from standard errors

```
> res <- predict(GLM, type="response",  
  newdata=Predictions, se.fit=TRUE)  
> transform(Predictions,  
  predicted=100*res$fit,  
  lwr=100*(res$fit - 1.96*res$se.fit),  
  upr=100*(res$fit + 1.96*res$se.fit))
```

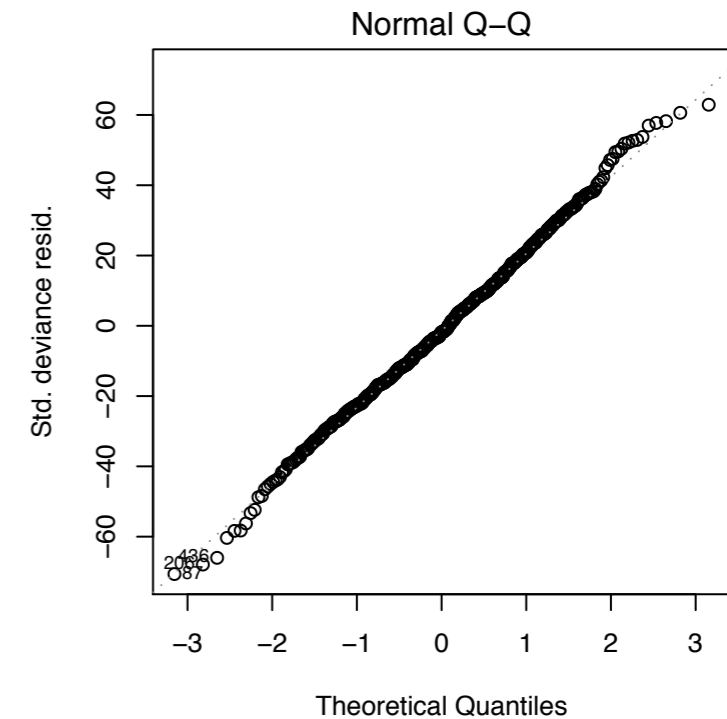
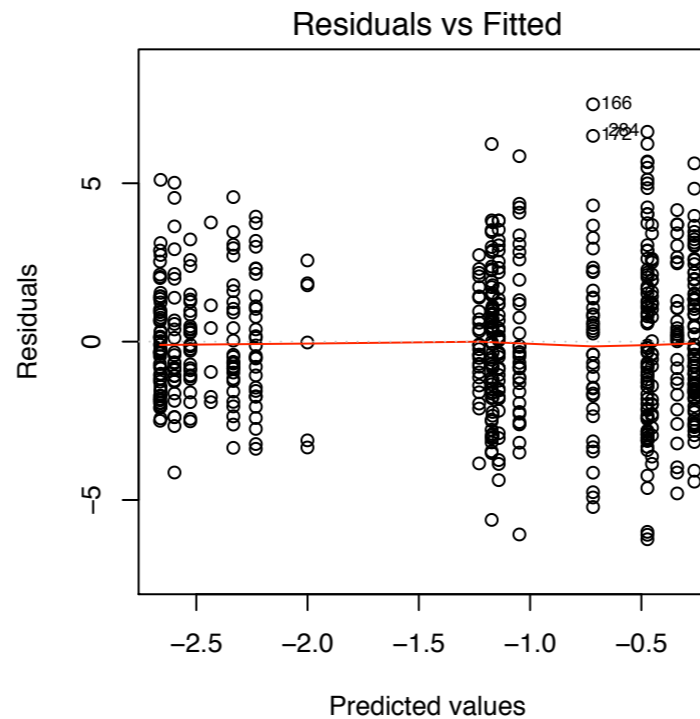
we can't compute prediction intervals for new texts — why?

Model diagnostics comparison

Linear Model



Generalised Linear Model



Still no satisfactory explanation for observed variation in frequency of passives between texts!

Take-home messages

- ◆ Don't trust statistic(ian)s blindly
 - You know how complex language really is!
 - linguists and statisticians should work together
- ◆ No excuse to avoid significance testing
 - good reasons to believe that binomial sampling distribution is a lower bound on variation in language
- ◆ Needed: large corpora with rich metadata
 - study & “explain” variation with statistical models
 - full data need to be available (not Web interfaces!)

T H A N K
Y O U

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