Introduction to Bayesian linear regression with brms — Part II: Bayesian Inference

Stefano Coretta 18/01/2020

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- 2. Frequentist inference.

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- 3. Bayesian inference:

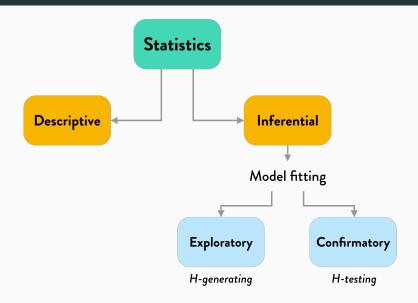
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STATISTICAL INFERENCE

Statistics



4

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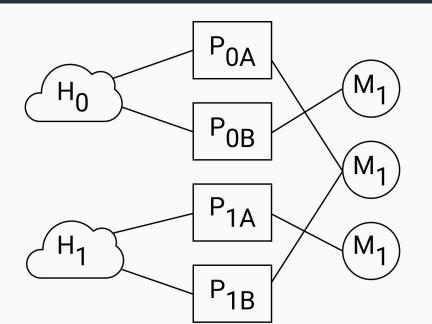
- Is there evidence for the hypothesis H?
- What is the strength of the evidence?

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 - It is important to decide in advance the details of the analysis.
 - Even when you think you are not making decisions, the model is.
- Inference is ultimately a long-term endeavour (via accumulation of knowledge).



FREQUENTIST INFERENCE

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- · Most of modern science is based on frequentism.
 - · lme4 package.
 - · Null Hypothesis Significance Testing.

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 - · Rejection of the Null Hypothesis (H_0) .
 - · No direct "support/evidence" for hypotheses.
- P-value (between 0 and 1).
 - Probability of obtaining an estimate as extreme or more extreme, assuming ${\cal H}_0$ is true.
 - · Should be as low as possible.

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- · No degrees of significance.

- · Significance is dichotomous.
 - \cdot p<lpha = "significant".
 - \cdot $p \geq \alpha$ = "non-significant".
- · No degrees of significance.
- "Significance" is a concept that makes sense only within frequentist statistics (NHST).

(Frequentist) confidence intervals

https://rpsychologist.com/d3/ci/

BAYESIAN INFERENCE

Bayesian inference

Bayesian statistics is based on the Bayesian interpretation of the Bayes theorem.

$$P(\theta \mid d) = \frac{P(d \mid \theta) \times P(\theta)}{P(d)}$$

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- · Strength of evidence.
- · Capitalise on previous knowledge.

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- · Inference from the posterior.
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- · Inference from the posterior.
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- · Inference using the Bayes factor.

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 - Condition B 95% CI = [-80, -15] ms.

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 - The posterior is compatible with H.

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- 1. Formulate a hypothesis:
 - H: Condition B decreases reaction times relative to Condition A by 100 ms.
- 2. Choose model specification (including priors).
- 3. Collect data.
- 4. Calculate the **posterior** (fit the model):
 - Condition B 95% CI = [-80, -15] ms.

- 1. Formulate a hypothesis:
 - H: Condition B decreases reaction times relative to Condition A by 100 ms.
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- 3. Collect data.
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- 2. Choose model specification (including priors).
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- 4. Calculate the **posterior** (fit the model):
 - Condition B 95% CI = [-80, -15] ms.
- 5. Inference:
 - The posterior suggests that Condition B decreases reaction times by 15 to 80 ms at 95% confidence.
 - The posterior is not compatible with H.

H₀ vs H₁

 H1 states that Condition B increases segment duration (alternative hypothesis), while H0 states that Condition B does not increase segment duration (null hypothesis, null effect).

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 - $\cdot H_1: \beta > 0$
 - $\cdot H_0: \beta = 0$

Region Of Practical Equivalence (ROPE):

. Define a region around $\beta=0$ that practically corresponds to a null effect.

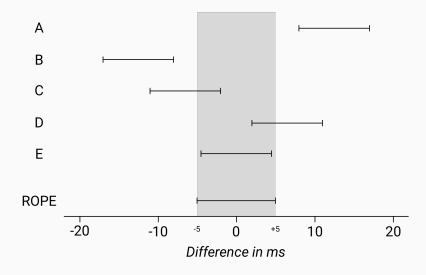
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 - · This ROPE has a width of 10 ms.
- Collect data until the 95% CI of β has a width equal to or smaller than the width of the ROPE.
 - Choose a minimal sample size (ideally based on a prospective power analysis).
 - Collect data and check 95% CI. If the width is greater than the ROPE, collect more data and repeat (sequential testing).



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- · Independent from the value of β .

Inference with a ROPE

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- Independent from the value of β .
- Higher precision means greater confidence in the estimated value of β .

The Bayes factor is the ratio of the likelihood of H1 to the likelihood of H2.

$$BF_{12}=\mathcal{L}(H_1)/\mathcal{L}(H_2)$$

BF p(M1 D) evidence 1-3 0.5-0.75 weak 3-20 0.75-0.95 positive 20-150 0.95-0.99 strong > 150 > 0.99 very strong			
3–20 0.75–0.95 positive 20–150 0.95–0.99 strong	BF	p(M1 D)	evidence
20–150 0.95–0.99 strong	1–3	0.5-0.75	weak
9	3-20	0.75-0.95	positive
> 150 > 0.99 very strong	20-150	0.95-0.99	strong
	> 150	> 0.99	very strong

```
priors <- c(
  prior(normal(0, 500), class = Intercept),
  prior(cauchy(0, 15), class = sigma),
  prior(normal(0, 750), class = b, coef = "vowelo"),
  prior(normal(0, 750), class = b, coef = "vowelu"),
  prior(cauchy(0, 15), class = sd),
  prior(lkj(2), class = cor)
)</pre>
```

```
f1_3_bf <- brm(
 f1 \sim 1 + vowel + (1 + vowel | speaker),
 family = gaussian(),
 prior = priors,
 data = f end,
 chains = 4,
 iter = 2000,
 file = "./cache/f1_3_bf",
 save_all_pars = TRUE
```

```
priors <- c(
  prior(normal(0, 500), class = Intercept),
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```

```
f1_3_place <- brm(
 f1 ~ 1 + vowel + c2_place + (1 + vowel + c2_place | s
 family = gaussian(),
 prior = priors,
 data = f end,
 chains = 4,
 iter = 2000,
 file = "./cache/f1_3_place",
 save all pars = TRUE
```

```
bf <- bayes factor(f1 3 bf, f1 3 place)</pre>
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
## Iteration: 6
## Iteration: 7
## Iteration: 8
## Iteration: 9
## Iteration: 1
## Iteration: 2
## Iteration: 3
```

bf

Estimated Bayes factor in favor of f1_3_bf over f1_3_place: 0.58852

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- BFs are VERY sensitive to priors and data.
- Always calculate and report BFs by comparing models with increasingly narrower priors (at least 3-4).
- It's important to run *sensitivity analyses* that assess the influece of the priors on the posterior (not only if you you BFs, but always).

THE END