Discussion: Classical p-values and the Bayesian posterior probability that the hypothesis is approximately true

Author: Brendan Kline Discussant: Ahnaf Rafi

Northwestern University

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### Summary I: high level stuff

- P-values are ubiquitous as a standard for what constitutes (sufficient) empirical evidence for scientific discovery.
- Usually framed in the context of rejecting an exact "neutral" null hypothesis in favor of ("more interesting") alternative(s).
- **This paper:** are p-values are actually informative about the (Bayesian posterior) probability that a null is **approximately correct**?
- Why? We construct tests for exact nulls, but draw conclusions about approximate nulls.

# Summary II: framing

- Throughout, for user-specified  $c, \varepsilon$ 
  - Exact null:  $H_0: \theta = c$ .
  - Approximate null:  $H_0^{\varepsilon}: \theta \in [c \varepsilon, c + \varepsilon]$
- Low p-value is associated (*conflated*) by researchers with "low probability" that  $\theta$  is "close to zero".
- Existing literature: If the prior has an atom at  $\theta = c$ , there is an increasing relationship between  $\Pr(\theta = c | \text{Data})$  and p-value.
- Paper's framing: in social sciences,
  - **()** Even if we test  $H_0$ , we really mean  $H_0^{\varepsilon}$  and draw conclusions about the latter.
  - 2 In addition, no reason to have prior with  $0 < \Pr(\theta = \tau) < 1$  for any  $\tau \in [c \varepsilon, c + \varepsilon]$ .

- **1** If we assume a continuous prior with positive density in  $[c \varepsilon, c + \varepsilon]$ , then p-value and  $\Pr(H_0^{\varepsilon}|\text{Data})$  no longer have an increasing relationship.
- 2 In particular,  $\Pr(H_0^{\varepsilon}|\text{Data})$  can be higher for lower p-values (and vice versa) suggests caution against using low p-values as a standard for judging empirical findings.
- Seven though main results are asymptotic, the phenomenon is true generally and in finite samples - not an "asymptotic curiosity".

I like the motivation of the particular Bayesian framework from what is done/said in practice:

- Bayesian approach is appropriate since we want to draw (probabilistic) conclusions about true values of the parameters.
- Approximate null and continuous priors are motivated by how researchers think about null hypotheses.
- The use of Bernstein-von Mises approximations is justified by the continuous prior.

### My take on broader implications of the paper

- Existing results on the increasing relationship between p-values and  $\Pr(H_0|\text{Data})$  with an atomic prior are internally consistent, but interpreting them outside their context is incorrect. The paper does a good job of driving home that point.
- Highlights the potential cost to the overall community of using p-values as a scientific standard: can miss out on treatment effects that are probably not close to zero.
- The large sample approximations provide one alternative standard for evaluating empirical findings.
- Provides a useful tool to retroactively assess evidence about whether published (non-zero) treatment effects are in fact likely close to zero.

#### Some criticisms

- The finite sample analysis is nice, but seems "tied" to the large sample results since sampling and posterior distributions are t and F distributions in finite samples.
- Some Monte-Carlo simulations with alternative continuous posteriors perhaps? (In lieu of closed form results.)
- Along those lines, asymptotic approximations are nice, but how good are they really? Is there something like a (uniform) Berry-Esseen bound for BvM?

## Controversial stuff (my own thoughts, time permitting) I

- To me, drawing conclusions about parameter values on the basis of p-values has always seemed awkward from a frequentist perspective:
  - Researchers can compute  $\Pr(\text{Data}|H_0)$ , but want to make statements about  $\Pr(H_0|\text{Data})$ .
  - By Bayes' theorem,

$$\begin{aligned} \Pr\left(H_{0}|\text{Data}\right) &= \frac{\Pr\left(\text{Data}|H_{0}\right)\Pr\left(H_{0}\right)}{\Pr\left(\text{Data}\right)} \\ &= \frac{\Pr\left(\text{Data}|H_{0}\right)\Pr\left(H_{0}\right)}{\Pr\left(\text{Data}|H_{0}\right)\Pr\left(H_{0}\right) + \Pr\left(\text{Data}|\neg H_{0}\right)\Pr\left(\neg H_{0}\right)}. \end{aligned}$$

Red = not available in frequentist world. Blue = basically what the p-value corresponds to.

• Additionally, there is a logical leap in the act of drawing conclusions about parameter values on the basis of p-values - they are computed "conditional" on both the **estimator** and the hypothesized value of the parameter.

# Controversial stuff (my own thoughts, time permitting) II

- As a standard for judging empirical findings, "small p-values" (Fisherian paradigm, null hypothesis significance testing [NHST]) thus seems rather strange.
- The Neyman-Pearson null+alternative, Type I + II error control paradigm is also not always helpful for evaluating scientific findings.
- Should not surprise anyone the p-value and confidence intervals are all about characterizing sampling error **assuming the null is true**.
- Not new, see e.g. Gigerenzer, Krauss, and Vitouch (2004), and Szucs and Ioannidis (2017) and references therein.
- Begs the question: from a frequentist perspective, how do we evaluate empirical findings without NHST?