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

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ABSTRACT. This paper relates *p*-values for the hypothesis that $\theta = c$ to the Bayesian posterior probability that the hypothesis is approximately true, in the sense that $\theta \in [c - \epsilon, c + \epsilon]$ for a selected $\epsilon > 0$. In a setup with a continuous prior for θ , the results show that a larger (respectively, smaller) *p*-value does not necessarily correspond to a higher (respectively, lower) probability that θ is close to *c*. Therefore, the results suggest caution about common ways of using *p*-values, specifically the use of small *p*-values as a key standard in empirical research.

Keywords: frequentist, hypothesis, posterior, testing

JEL classification: C11, C12, C18

1. INTRODUCTION

This paper is about *p*-values for the hypothesis $\theta = c$, where θ is a scalar parameter and *c* is a known constant (e.g., often c = 0). An important question is how to use *p*-values to draw conclusions about θ . For example, one common usage of *p*-values concerns the case where θ is a treatment effect, like a regression coefficient. In that case, in practice to establish and publish an empirical finding that the treatment has an effect on an outcome, it is considered important to find a small *p*-value for the hypothesis that the treatment effect is zero. This is a running example, but the results hold regardless of the meaning of θ .

The results of Edwards, Lindman, and Savage (1963), Berger and Delampady (1987), Berger and Sellke (1987), Sellke, Bayarri, and Berger (2001), Held and Ott (2016), and Kline

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(2022), as representative of a related literature, establish that it is possible to "calibrate" a p-value via the corresponding minimum posterior probability that $\theta = c$. In the setup of that literature, there is a positive prior probability that $\theta = c$, and a *class* of priors for θ on $\theta \neq c$. The "minimum" is taken over this class of priors. Although the *p*-value is (much) smaller than the minimum posterior probability, there is a monotone increasing relationship between the *p*-value and the minimum posterior probability of the hypothesis that $\theta = c.^{12}$ Essentially, this literature implies that only very small p-values are enough to "reject" that $\theta = c$. Conversely, larger *p*-values correspond to relatively higher minimum posterior probabilities that $\theta = c$. This provides a Bayesian justification for "rejecting" the hypothesis based on a sufficiently small *p*-value, and "not rejecting" the hypothesis based on larger *p*-values. Put differently, this provides a justification for using small *p*-values as a standard for empirical evidence that θ is unlikely to be equal to c. Indeed, the Benjamin, Berger, Johannesson, Nosek, Wagenmakers, et al. (2018) proposal to redefine "statistical significance" to be a *p*-value smaller than 0.005 is based in part on considerations of how *p*-values relate to the (minimum) posterior probability that $\theta = c$. Also, of course, with classical hypothesis testing without any reference to Bayesian analysis, smaller *p*-values are interpreted as rejecting the hypothesis, and larger p-values lead to not rejecting the hypothesis and/or the conclusion that there is no empirical evidence against the hypothesis.³

In practice, often the empirical question is not whether $\theta = c$ is literally true but rather whether θ is "close" to c. Specifically, consider " ϵ -approximate" hypotheses of the form

¹There are many "minimum posterior probabilities" proposed in that literature, based on different classes of priors over which the minimum is taken, and monotonicity is a shared property.

²Lindley's paradox (e.g., Jeffreys (1939); Lindley (1957)) establishes conditions for the *p*-value to be less than the posterior probability of the hypothesis for a (uniform) single prior on $\theta \neq c$. A review of the literature on minimum posterior probabilities can be found in Held and Ott (2018).

³Along similar lines to the case of not rejecting the hypothesis, using a "limited information" posterior that conditions on the dichotomous event of rejecting (or not) the hypothesis $\theta = c$ by classical methods, Abadie (2020) shows the posterior conditional on not rejecting the hypothesis has much higher probability mass at $\theta = c$ compared to the prior, whereas the posterior conditional on rejecting the hypothesis is not too different from the prior.

 $\theta \in [c - \epsilon, c + \epsilon]$ for some specified $\epsilon > 0$ chosen by the econometrician.⁴ An ϵ -approximate hypothesis is about θ being "close" to c. In the setup with a positive prior probability that $\theta = c$, it can also be justified to use small p-values as a standard for empirical evidence that θ is unlikely to be "close" to c, using the same "calibration" of p-values for the hypothesis $\theta = c$. The reason: when there is positive probability that $\theta = c$ and ϵ is small enough, the posterior probability of $\theta = c$ is approximately the same as the posterior probability of $\theta \in [c - \epsilon, c + \epsilon]$. See Remark 4 for more on this point.

However, particularly in the social sciences, often there is zero prior probability that $\theta = c$. So, in that setting, how does the *p*-value relate to the posterior probability that θ is close to c? In other words, in that setting, is there a (Bayesian) justification for using small *p*-values as a key standard for empirical evidence?⁵ This paper relates *p*-values for the hypothesis that $\theta = c$ to the Bayesian posterior probability of $\theta \in [c - \epsilon, c + \epsilon]$ in a setup with a continuous prior, with zero prior probability that $\theta = c$.

The results allow the econometrician to choose any ϵ . ϵ can be a selected constant or data-dependent. It is possible to interpret ϵ as representing the magnitude of deviations from $\theta = c$ that are viewed as "economically significant." In particular, because ϵ can be datadependent, in the case that θ reflects a treatment effect, ϵ can be taken to be proportionate to the (estimated) effect of a reference treatment, so that the ϵ -approximate hypothesis relates to the effect of the treatment under study relative to the reference treatment. Although "economic significance" is the most plausible determining factor for ϵ , other considerations are possible; in particular, ϵ is allowed to be data-dependent, which allows for a variety of specifications. Similar to other hypothesis testing problems, the scaling of the model parameters is an important consideration; θ , c, and ϵ are all defined in terms of the same scale (units of measurement).

⁴It does not matter if the ϵ -approximate hypothesis is defined as a closed interval or open interval.

⁵In that setting it is not interesting to ask about the posterior probability that $\theta = c$ exactly, since that is necessarily zero.

p-VALUES AND APPROXIMATE HYPOTHESES

The paper derives a closed-form expression for the posterior probability of the ϵ -approximate hypothesis, as a function of the *p*-value (and classical estimate of θ). The main results are large sample approximations, which are valid quite generally in parametric and semiparametric models when the Bernstein-von Mises phenomenon holds. There are also finite sample results, based on distributional assumptions on the data generating process and a uniform prior. The large sample and finite sample results are very similar. Consequently, the results are not tied specifically to a large sample approximation or finite sample analysis. Then, it is possible to investigate the relationship between the *p*-value and the posterior probability of the ϵ -approximate hypothesis.

As a point about the positioning of this paper, it is important to note that this paper follows the tradition in econometrics of exploring the properties of the sorts of empirical methods used in practice and the interpretations that can be applied to them.⁶ Here the analysis focuses on the *p*-value, and interprets and relates that to the ϵ -approximate hypotheses. The point of this sort of analysis is to investigate the properties of the empirical methods used in practice. As discussed below, the results have important practical implications, given the use of *p*-values. See also Remark 4 for more discussion of possible alternative empirical analyses.

In short, the results of this paper show that a small *p*-value is neither necessary nor sufficient for there to be a small posterior probability that θ is close to *c*. In particular, relatively larger *p*-values can correspond to relatively smaller posterior probabilities that θ is close to *c*.

This has important implications for using and interpreting *p*-values. As a running example, if θ is a treatment effect (e.g., a regression coefficient), the results can be used to relate a *p*-value for the hypothesis $\theta = 0$ to the probability that the treatment effect is close to 0.

⁶In a general way, this is similar to how the local average treatment effects literature (e.g., Imbens and Angrist (1994)) or the recent literature on two-way fixed effects (and related) estimators (e.g., De Chaisemartin and d'Haultfoeuille (2020), Borusyak, Jaravel, and Spiess (2021), Callaway and Sant'Anna (2021), Goodman-Bacon (2021), Sun and Abraham (2021), and Athey and Imbens (2022), among others) investigates the meaning of standard IV estimators or standard TWFE estimators, and how they can be interpreted in relationship to treatment effect heterogeneity. Because these estimators (like IV or TWFE estimators) are often used in settings involving treatment effect heterogeneity, an important question concerns how the standard estimators relate to treatment effect heterogeneity.

Concretely, close to 0 would mean that $\theta \in [-\epsilon, \epsilon]$ for $\epsilon > 0$ chosen by the econometrician. In this treatment effects setting, the results suggest caution about the practice of using small *p*-values as a key standard for empirical evidence, like the standard for finding/publishing an effect of a treatment on an outcome.⁷ In particular the results show that a larger (respectively, smaller) *p*-value for the hypothesis that the treatment effect is zero does not necessarily correspond to a higher (respectively, lower) probability that the treatment effect is close to zero. Comparing two studies, a study reporting a larger *p*-value can actually have a smaller posterior probability of a treatment effect that is close to zero. This suggests caution against using small *p*-values as a key standard for empirical evidence, as in cutoff rules for determining "significance." This contrasts with the previously discussed results from the literature that do provide justifications for using small *p*-values as a standard in different settings/setups.

Of particular importance, the results imply the research community can "miss" (e.g., not publish) treatment effects that are probably *not* close to zero if the research community only "accepts" (or focuses on) treatment effects with small *p*-values for the hypothesis of zero treatment effect. Formally, "probably not close to zero" refers to (one minus) the probability of the ϵ -approximate hypothesis. The expressions derived in this paper make it possible to use a *p*-value to assess the posterior probability of the ϵ -approximate hypothesis. This can be viewed as either a complement or substitute to the standard approach of reporting *p*-values in empirical research; in particular, it is possible to use these expressions to assess the posterior probability of the ϵ -approximate hypothesis from previous research even without access to the original data, in the sense that the expressions depend only on quantities that are conventionally reported in research.

These results contribute to the broader discussion surrounding the use of p-values, including a statement by the American Statistical Association in Wasserstein and Lazar (2016), comment

⁷It is clear that standards for empirical research are such that a small *p*-value is important. Some evidence is that Masicampo and Lalande (2012), Leggett, Thomas, Loetscher, and Nicholls (2013), and Brodeur, Lé, Sangnier, and Zylberberg (2016) find high prevalence of *p*-values around and just below the "significance cutoff" of 0.05.

in Nature in Amrhein, Greenland, and McShane (2019), and various statements on *p*-values at specific journals or for specific fields (e.g., Trafimow and Marks (2015), Harvey (2017), Gill (2018), Harrington, D'Agostino Sr, Gatsonis, Hogan, Hunter, Normand, Drazen, and Hamel (2019)). The results also contribute to related literatures that relate classical inference and Bayesian inference in other ways, including Liao and Jiang (2010), Kline (2011), Moon and Schorfheide (2012), Kline and Tamer (2016), Chen, Christensen, and Tamer (2018), Gafarov, Meier, and Montiel Olea (2018), Liao and Simoni (2019), and Giacomini and Kitagawa (2021) for partially identified models and Chernozhukov and Hong (2003), Kitagawa, Montiel Olea, Payne, and Velez (2020), and Liu, Li, Yu, and Zeng (2022) for relating classical inference results to (quasi) Bayesian approaches.

1.1. Notation. Notation is standard. The $m \times m$ identity matrix is $I_{m \times m}$. The L^p norm of $x \in \mathbb{R}^m$ is $||x||_p$. Let $\delta_{\Sigma,2}(x, y) = ||\Sigma^{-\frac{1}{2}}(x-y)||_2$ for a positive definite Σ . The total variation distance between random variables X and Y is $||X - Y||_{TV}$. For a sequence $X_N, X_N \to^p X$ means converges in probability to X and $X_N \to^d X$ means converges in distribution to X. The indicator 1[E] of logical statement E is 1 when E is true and 0 otherwise. As convention, z1[false] = 0; for example, $\frac{x}{y}1[y \neq 0] = 0$ when y = 0. A multivariate normal distribution with mean μ and covariance Σ is $\mathcal{N}(\mu, \Sigma)$. A (central) chi-squared distribution with m degrees of freedom is χ^2_m . A noncentral chi-squared distribution with m degrees of freedom and non-centrality parameter λ is $\chi^2_{m,\lambda}$.⁸ For a random variable X, the cumulative distribution function is $F_X(\cdot)$, the quantile function is $Q_X(\cdot)$, the complementary cumulative distribution function is $\overline{F}_X(\cdot) \equiv 1 - F_X(\cdot)$, and $X(\mathcal{B}) \equiv P(X \in \mathcal{B})$.

2. Setup

The data is a sample of N i.i.d. observations from P_0 , the true data generating process, so the data is $X^{(N)} \equiv \{X_i\}_{i=1}^N$ where $X_i \sim^{iid} P_0$. There also is a statistical model, where $\overline{{}^8\text{This paper uses the parameterization where } E(\chi^2_{m,\lambda}) = m + \lambda.$ $\psi = (\theta, \gamma_2, \dots, \gamma_t, \dots, \gamma_m)$ is the finite-dimensional parameter of the model, with parameter space $\Psi \subseteq \mathbb{R}^m$ for some m. The true value of ψ is ψ_0 . The *p*-values and posterior probabilities will concern the scalar parameter of interest θ .

The "statistical model" can be a fully parametric model or a semi-parametric model. Specifically, per Remark 1 about semi-parametric models, there can be infinite-dimensional nuisance parameters. Overall, the analysis proceeds by making a high-level assumption on the classical estimator and posterior distribution, which is known to hold across a variety of types of models. As a consequence, the setup does not need to specify the use of any particular statistical model. As a further consequence, the analysis applies both to parametric and semi-parametric models, and in either case, the analysis applies to the scalar parameter of interest θ .

Specifically, the analysis presumes there is a classical ("frequentist") estimator $\hat{\psi}_N = (\hat{\theta}_N, \hat{\gamma}_{N,2}, \dots, \hat{\gamma}_{N,t}, \dots, \hat{\gamma}_{N,m})$ of ψ , which is the basis for the *p*-value. Also there is a Bayesian posterior for ψ , denoted $\Pi_{\psi|X^{(N)}}(\cdot)$ so $\Pi_{\psi|X^{(N)}}(A)$ is the posterior probability that $\psi \in A$ conditional on the data. Sometimes, rather than $\Pi_{\psi|X^{(N)}}(A)$, the equivalent notation $\Pi(\psi \in A|X^{(N)})$ is used.

It is worth observing that the posterior in this paper conditions on the entire dataset, so this paper relates *p*-values to a conventional Bayesian analysis based on the entire dataset. A potential alternative analysis would analyze a "non-standard" posterior that conditions on "less" information, as discussed for instance in Footnote 3. Although reasonable, by construction this would not correspond to the "standard" posterior the Bayesian would get from analyzing the data in the "standard" way, and as such, that approach is not taken here.

3. Large sample approximation results

This section establishes the large sample approximation results that relate the *p*-value for the hypothesis that $\theta = c$ to $\Pi(\theta \in [c - \epsilon, c + \epsilon]|X^{(N)})$. The latter is the Bayesian posterior probability that the hypothesis is approximately true. Intuitively, the large sample approximation established in this section is a function $\pi_{c,\epsilon}(p_N,\hat{\theta}_N)$, that depends on the *p*-value and the classical estimate $\hat{\theta}_N$, such that $\Pi(\theta \in [c - \epsilon, c + \epsilon] | X^{(N)}) \approx \pi_{c,\epsilon}(p_N, \hat{\theta}_N)$ in large samples. Both $\Pi(\theta \in [c - \epsilon, c + \epsilon] | X^{(N)})$ and $\pi_{c,\epsilon}(p_N, \hat{\theta}_N)$ are functions of $X^{(N)}$.

More formally, the large sample approximation establishes that, for any given $\delta > 0$, the P_0 probability of the set of realizations $X^{(N)}$ such that $\left|\Pi(\theta \in [c - \epsilon, c + \epsilon]|X^{(N)}) - \pi_{c,\epsilon}(p_N, \hat{\theta}_N)\right| < \delta$ approaches one as sample size increases. In other words, $\Pi(\theta \in [c - \epsilon, c + \epsilon]|X^{(N)}) \approx \pi_{c,\epsilon}(p_N, \hat{\theta}_N)$ for all but an exceptional set of realizations $X^{(N)}$, and the P_0 -probability of the exceptional set decreases to zero as sample size increases.

This large sample approximation is a statement that applies to realizations $X^{(N)}$ from P_0 , assuming that $X^{(N)}$ is an i.i.d. sample from P_0 , as is the case with other large sample approximations of Bayesian posteriors (e.g., Van der Vaart (1998, Chapter 10)). The P_0 distribution is used both as the underlying data generating process, and to "measure" the set of realizations $X^{(N)}$ where the approximation holds. Although the derivation of the *p*-value involves repeating sampling considerations, from the point of view of the large sample approximation of the Bayesian posterior, the *p*-value is simply a particular function of the realization $X^{(N)}$ in the same way that the Bayesian posterior is a particular function of the realization $X^{(N)}$. Large sample approximations of Bayesian posteriors can be distinguished from the classical repeating sampling approximations used in frequentist analysis, as those concern the *distribution* of an estimator that is induced by the sampling process.

The large sample approximation results are based on an assumption about the relationship between the sampling distribution of $\hat{\psi}_N$ and the posterior distribution for ψ .

Assumption 1. It holds that:

- (1) The classical estimator $\hat{\psi}_N$ is asymptotically normal, in the sense that $\sqrt{N}(\hat{\psi}_N \psi_0) \rightarrow^d \mathcal{N}(0, \Sigma_0)$ where Σ_0 is nonsingular. There is a consistent estimator $\hat{\Sigma}_N$ of Σ_0 .
- (2) The Bayesian posterior for ψ is asymptotically normal, in the sense that $||\Pi_{\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} \mathcal{N}(0,\Sigma_0)||_{TV} \to^p 0.$

The analysis treats Σ_0 as an unknown quantity, with corresponding estimate $\hat{\Sigma}_N$. A possible alternative analysis is the known Σ_0 case. Because Σ_0 typically depends on unknown model parameters, it is unlikely that Σ_0 would be known in empirical practice, so it must be estimated using the data. Further, the known Σ_0 case is not particularly interesting from the perspective of this paper: If Σ_0 were known, then $\hat{\theta}_N$ (implicitly along with N and c) would be enough information to compute the *p*-value, and therefore reporting a *p*-value would convey no additional information beyond the information from reporting $\hat{\theta}_N$. But, when Σ_0 is unknown, as in empirical practice, the *p*-value does contain additional information beyond $\hat{\theta}_N$, which justifies the empirical practice of reporting *p*-values in addition to $\hat{\theta}_N$. Indeed, a key driving force of the results is that the p-value contains information about Σ_0 . This is relevant information since the posterior distribution depends on Σ_0 . This condition on the covariance corresponds also to the analogous condition on the posterior used in the finite sample results in Section 6. This provides another intuition for why this condition on the covariance is used; in the finite sample results, it arises because of the unknown (co)variance of the underlying data generating process. Correspondingly, essentially the same condition also holds "in the limit" in the large sample approximation.

Assumption 1 holds as the consequence of theorems commonly known as "Bernstein-von Mises theorem(s)." The following remark quickly summarizes settings where the "Bernstein-von Mises theorem(s)" hold. The results of this section apply to those settings.

Remark 1 (Sufficient conditions for Assumption 1). Assumption 1 holds in these settings:

(1) **Parametric model**: Consider a parametric model P_{ψ} with finite-dimensional parameter ψ . Then $P_0 = P_{\psi_0}$, where ψ_0 is the true value. Often, $\hat{\psi}_N$ is the maximum likelihood estimate of ψ , and Σ_0 the inverse Fisher information matrix. The prior for ψ must have a continuous and positive density on a neighborhood of ψ_0 . Under a few further regularity conditions, the parametric Bernstein-von Mises results (e.g., Le Cam (1986, Chapter 12), Van der Vaart (1998, Chapter 10), Le Cam and Yang

(2000, Chapter 8)) imply Assumption 1 is satisfied. The regularity conditions are similar to those used in the familiar approaches to asymptotic normality of parametric maximum likelihood estimation, see for instance the discussion in Van der Vaart (1998, Chapter 10). Although there are different ways to state these regularity conditions, they generally involve identifiability of the model, differentiability in quadratic mean or local asymptotic normality, and a separation condition which itself can be shown to follow from regularity conditions like compactness of the parameter space and continuity of P_{ψ} as a function of ψ .

(2) Semi-parametric model: Consider a semi-parametric model $P_{\psi,\eta}$ with finitedimensional parameter ψ and infinite-dimensional parameter η . Then $P_0 = P_{\psi_0,\eta_0}$, where (ψ_0,η_0) is the true value. Often, $\hat{\psi}_N$ can be taken to be an asymptotically linear and efficient estimator of ψ , and Σ_0 the inverse efficient Fisher information matrix. The marginal prior for ψ must have a continuous and positive density on a neighborhood of ψ_0 . Under a few further regularity conditions, the semi-parametric Bernstein-von Mises results (e.g., Shen (2002), Bickel and Kleijn (2012), Castillo (2012), Castillo and Rousseau (2015)) imply Assumption 1 is satisfied. The regularity conditions generally are semi-parametric parallels to the regularity assumptions mentioned above. See for instance the discussion in Rousseau (2016).

Overall, the results in this section are large sample approximations, for any setup compatible with Assumption 1. Remark 1 is not a comprehensive review. There are Bernstein-von Mises results that cover certain nonparametric models (e.g., Castillo and Nickl (2013)), and important specific models, including limited information and moment condition models (e.g., Kwan (1999), Kim (2002), and Chib, Shin, and Simoni (2018)), linear and partially linear regressions (e.g., Bickel and Kleijn (2012) and Norets (2015)) proportional hazard models (e.g., Kim (2006)), and models with quasi-posteriors (e.g., Chernozhukov and Hong (2003)).

Assumption 1 may not hold if the model is misspecified as in Kleijn and Van der Vaart (2012) and Müller (2013).

Remark 2 (Parameter of interest is a function of ψ). Suppose Assumption 1 holds, but the parameter of interest is $\tilde{\psi} \in \mathbb{R}^1$ with $\tilde{\psi} = f(\psi)$ for some known function $f(\cdot) : \mathbb{R}^m \to \mathbb{R}^1$ that is continuously differentiable at ψ_0 . Suppose the $(1 \times m)$ -matrix of first derivatives of $f(\cdot)$ is $F(\cdot)$, and $F(\psi_0)$ is non-zero. The delta theorem (e.g., Van der Vaart (1998, Chapter 3), Wasserman (2004)) implies that Assumption 1 holds for $\tilde{\psi}$, with $\hat{\psi}_N$ replaced by $f(\hat{\psi}_N)$, ψ_0 replaced by $f(\psi_0)$, ψ replaced by $f(\psi)$, and Σ_0 replaced by $F(\psi_0)\Sigma_0F(\psi_0)'$. Therefore, the results apply to a parameter of interest that is a continuously differentiable function of a parameter that satisfies Assumption 1.

The Wald test is the classical method for testing the hypothesis $\theta = c$. The Wald test statistic is

$$W_N = N(\hat{\theta}_N - c)'(\hat{\Sigma}_{N,11})^{-1}(\hat{\theta}_N - c).$$

By Assumption 1, if the hypothesis $\theta = c$ is true, W_N converges in distribution to χ_1^2 . Therefore, the *p*-value for the hypothesis that $\theta = c$ is

$$p_N = P(\chi_1^2 > W_N) = \overline{F}_{\chi_1^2}(W_N).$$

Theorem 1. Under Assumption 1, and when the p-value is not 1,⁹ for any $\epsilon > 0$ chosen by the econometrician (possibly data-dependent per Remark 5), the Bayesian posterior probability of the ϵ -approximate hypothesis that $\theta \in [c - \epsilon, c + \epsilon]$ is approximately $F_{\chi^2_{1,Q_{\chi^2_1}(1-p_N)}}\left(\frac{\epsilon^2}{|\hat{\theta}_N - c|^2}Q_{\chi^2_1}(1-p_N)\right)$ in large samples. Formally, $\left(\prod\left(\theta \in [c - \epsilon, c + \epsilon]|X^{(N)}\right) - F_{\chi^2_{1,Q_{\chi^2_1}(1-p_N)}}\left(\frac{\epsilon^2}{|\hat{\theta}_N - c|^2}Q_{\chi^2_1}(1-p_N)\right)\right)1[p_N < 1] \rightarrow^p 0.$

 $^{{}^{9}}p_{N} = 1$ exactly happens with asymptotic probability 0, and finite sample probability 0 with a continuous sampling distribution, so this "exclusion" is irrelevant formally in the asymptotic approximation and also for all practical purposes. Relatedly, $p_{N} = 0$ exactly cannot happen.

All proofs are collected in Appendix A, part of the main paper. Theorem 1 relates the *p*-value to the posterior probability of the ϵ -approximate hypothesis. Figure 1 displays the contour plot of the posterior probability of the ϵ -approximate hypothesis as a function of the *p*-value and $\frac{|\hat{\theta}_N - c|}{\epsilon}$. The latter quantity depends on ϵ . To illustrate by example, if $\frac{|\hat{\theta}_N - c|}{\epsilon} = 1.5$ and the *p*-value is 0.05, Figure 1 indicates the posterior probability of the ϵ -approximate hypothesis is 0.2562. Note that because Σ_0 is unknown (and therefore must be estimated), $\hat{\theta}_N$ (and *c* and ϵ) does not uniquely determine the *p*-value. The figures in this paper have axes that are truncated away from *p*-values of 0 and 1 and away from values of $\frac{|\hat{\theta}_N - c|}{\epsilon}$ of 0 for the same reason as described in Footnote 9. This has no meaningful impact on the interpretation of the figures. For values outside this range, the formula in the theorem can be used. For a different perspective, Figure 2 displays the posterior probability of the ϵ -approximate hypothesis as a function of the *p*-value, for various values of $\frac{|\hat{\theta}_N - c|}{\epsilon}$. Some of the qualitative features revealed in Figure 2 have corresponding theoretical results in Corollary 1 and Corollary 2.

It may be tempting to think that somehow this result is an asymptotic curiosity, but Section 6 shows essentially the same result obtains also in a finite sample analysis. Therefore, the main point of this paper is not tied to a particular setup (e.g., "asymptotic approximation" or "finite sample") but rather a general feature of *p*-values. The analysis of the large sample approximation reduces the dependence on a particular model setup or prior distribution, but the large sample approximation is not the "reason" for the result, since essentially the same result arises in the finite sample analysis. Along similar lines, it is worth noting that the large sample approximation in Theorem 1 and Figure 1 exists as a non-degenerate quantity as a direct implication of the non-degenerate large sample approximation in Assumption 1. Of course, per consistency of the posterior distribution, the posterior distribution of $\sqrt{N}(\psi - \hat{\psi}_N)$ does not converge to a point mass, per Assumption 1. Using the large sample approximation to the posterior distribution of $\sqrt{N}(\psi - \hat{\psi}_N)$, it is possible to have non-degenerate large



FIGURE 1. Posterior probability of the ϵ -approximate hypothesis that $\theta \in [c - \epsilon, c+\epsilon]$ for given *p*-value and $\frac{|\hat{\theta}_N - c|}{\epsilon}$ according to the large sample approximation.

sample approximations to the posterior distribution of ψ . By way of comparison to a more familiar situation with sampling distributions, this is analogous to how the large sample approximation to the sampling distribution of $\sqrt{N}(\hat{\psi}_N - \psi_0)$ is non-degenerate, making it possible to have non-degenerate large sample approximations to the sampling distribution of $\hat{\psi}_N$, which in particular form the basis of standard approaches to hypothesis testing. In particular, it may be worth noting that the formula for the *p*-value depends on sample size, and therefore the large sample approximation derived here does indeed involve the typical "scaling" by sample size, since the large sample approximations are functions of the *p*-value that depends on sample size.



FIGURE 2. Posterior probability of the ϵ -approximate hypothesis that $\theta \in [c - \epsilon, c + \epsilon]$ as function of *p*-value, for various $\frac{|\hat{\theta}_N - c|}{\epsilon}$, according to the large sample approximation.

Shortly, Section 4 discusses the main implications of this result for using p-values in practice. Before getting there, it is worthwhile to discuss a few intermediate implications of this result.

One implication of this result is that it shows the posterior probability of the ϵ -approximate hypothesis depends on $\hat{\theta}_N$. The *p*-value by itself is not determinative of the posterior probability of the ϵ -approximate hypothesis. This contrasts with the fact that the *p*-value by itself is used to decide on rejecting or not the hypothesis that $\theta = c$ by classical standards, or other "cutoff rules" for significance like Benjamin, Berger, Johannesson, Nosek, Wagenmakers, et al. (2018), and the fact that the minimum posterior probability of the hypothesis that $\theta = c$ discussed in the introduction depends only on the *p*-value (and also the discussion of a related relationship for the ϵ -approximate hypothesis in that setup).

Another implication of this result concerns the relationship between the posterior probability of the ϵ -approximate hypothesis and the *p*-value.

First, consider the case that the classical estimate provides "evidence against" the ϵ -approximate hypothesis, in the sense that $\hat{\theta}_N \notin [c - \epsilon, c + \epsilon]$, equivalent to $\frac{|\hat{\theta}_N - c|}{\epsilon} > 1$. Corollary 1 shows in that case that there is a non-monotone relationship between the *p*-value and the posterior probability of the ϵ -approximate hypothesis, even for given $\hat{\theta}_N$. This can be seen in Figure 1 and Figure 2. Both small *p*-values and large *p*-values correspond to a small posterior probability of the ϵ -approximate hypothesis. This contrasts with the fact that small *p*-values lead to rejecting the hypothesis that $\theta = c$ and large *p*-values do not, whether by classical standards or other "cutoff rules" for significance, and the fact that there is a monotone relationship between the *p*-value and the minimum posterior probability of the hypothesis that $\theta = c$. A large *p*-value can mean that the ϵ -approximate hypothesis is unlikely to be true, because values of the parameter other than the ϵ -approximate hypothesis values are more likely. Essentially, this effect is driven by what the *p*-value says about the variance of the posterior, which in turn affects the posterior probability of the ϵ -approximate hypothesis.

Corollary 1. Under the same conditions as Theorem 1, for fixed $\hat{\theta}_N$ with $\frac{|\hat{\theta}_N - c|}{\epsilon} > 1$: The asymptotic approximation to the posterior probability that $\theta \in [c - \epsilon, c + \epsilon]$ is maximal (as a function of p_N) when $p_N = p_{N,max}$, where $p_{N,max} = \overline{F}_{\chi_1^2} \left(\frac{1}{2\sqrt{\epsilon_N}} \log\left(\frac{-\sqrt{\epsilon_N}-1}{\sqrt{\epsilon_N}-1}\right)\right)$ with $\tilde{\epsilon}_N = \frac{\epsilon^2}{|\hat{\theta}_N - c|^2}$. The asymptotic approximation to the posterior probability of the ϵ -approximate hypothesis is an increasing function of p_N when $p_N \in (0, p_{N,max})$ and a decreasing function of p_N when $p_N \in (p_{N,max}, 1)$. As $p_N \to 0$ or $p_N \to 1$, the asymptotic approximation to the posterior probability of the ϵ -approximate hypothesis limits to 0.

Second, now consider the case that the classical estimate provides "evidence for" the ϵ -approximate hypothesis, in the sense that $\hat{\theta}_N \in [c - \epsilon, c + \epsilon]$, equivalent to $\frac{|\hat{\theta}_N - c|}{\epsilon} \leq 1.^{10}$ Corollary 2 shows in that case that the probability of the ϵ -approximate hypothesis is greatest when the *p*-value is smallest, for given $\hat{\theta}_N$. This can be easily seen in Figure 1 and Figure 2. Obviously, this is the exact opposite of common practice, where small *p*-values are used to reject the hypothesis.

Corollary 2. Under the same conditions as Theorem 1, for fixed $\hat{\theta}_N$ with $0 < \frac{|\hat{\theta}_N - c|}{\epsilon} \leq 1$: The asymptotic approximation to the posterior probability that $\theta \in [c - \epsilon, c + \epsilon]$ is a decreasing function of p_N . As $p_N \to 1$, the asymptotic approximation to the posterior probability of the ϵ -approximate hypothesis limits to 0. As $p_N \to 0$, the asymptotic approximation to the posterior to the posterior probability of the ϵ -approximate hypothesis limits to 0. As $p_N \to 0$, the asymptotic approximation to the case the limit is $\frac{|\hat{\theta}_N - c|}{\epsilon} = 1$ in which case the limit is $\frac{1}{2}$.

The general question of the relative merits of classical inference and Bayesian inference is substantially beyond the scope of this paper, but Efron (1986) suggests overall that there are "powerful theoretical reasons for preferring Bayesian inference." One such argument concerns the likelihood principle (e.g., Birnbaum (1962)) and its perceived support for Bayesian inference (e.g., Berger and Wolpert (1988)). Another such argument concerns decision theory and subjective probability foundations (e.g., de Finetti (1970, 1975); Savage (1954, 1972)). Such arguments in favor of Bayesian inference are the underpinning of the motivation for studying the question considered in this paper, as well as the related literature cited in the introduction, as they provide the basic justification for wanting to know what a Bayesian inference would conclude. A researcher who is even partly convinced by the arguments in favor of Bayesian inference can find it important to know how the Bayesian conclusion differs

¹⁰Again, per Theorem 1, the (asymptotically) probability 0 - and thus irrelevant for practical purposes - event that $\hat{\theta}_N = c$ exactly is excluded from the results/discussion.

¹¹This is a bit difficult to see in Figure 1 since the posterior probability is "close" to 1 only for *extremely* small p_N , for some $\frac{|\hat{\theta}_N - c|}{\epsilon}$.

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from the classical conclusion. Of course, Bayesian inference can also be criticized, for example along the lines outlined by Efron (1986), which include concerns about implementing Bayesian inference in practice and concerns about the "objectivity" of Bayesian inference due to the use of a prior distribution. These concerns are contextual to any particular application of Bayesian inference. And, as it turns out, those concerns apply in only a somewhat limited degree to the analysis done in this paper, since this paper indeed does do the Bayesian inference (addressing the concern about implementation in practice) and since this paper uses a large sample approximation (which sidesteps the issues about the prior distribution, as discussed above).

4. Implications of the results for the use of p-values in practice

The results suggest caution against the use of small *p*-values as a key standard in empirical research, including via *p*-value cutoff rules for "significance." The results of this paper show that a small *p*-value is neither necessary nor sufficient for there to be a small posterior probability that θ is close to *c*. On the other hand, it is possible to use the expressions derived in this paper to assess the posterior probability that θ is close to *c*. This can be done even without access to the original data, since the expressions can be applied to a reported *p*-value and classical estimate.

To illustrate by a numerical example, suppose θ is the effect of a treatment on an outcome, like a regression coefficient, and the *p*-value is for the hypothesis $\theta = 0$. For this example, a "close to zero" effect is any effect in [-0.1, 0.1], so $\epsilon = 0.1$. Nothing changes if θ and ϵ are scaled by the same positive constant. Consider these specifications for the *p*-value and classical estimate of the effect: $(p = 0.04, \hat{\theta} = 0.12)$ and $(p = 0.11, \hat{\theta} = 0.14)$. The first specification has a smaller *p*-value, which has the familiar implications: it implies the first specification has a smaller minimum posterior probability that the effect is zero, and it implies rejecting the hypothesis that the effect is zero by classical standards for the first specification. The *p*-value for the second specification does not achieve "statistical significance" and so it might be difficult to "sell"/publish such a result. It might be said that the second specification shows a "lack of evidence" against the hypothesis that the treatment effect is (close to) zero. Yet, the first specification implies a 36.6% posterior probability of an effect that is close to zero, greater than the 32.1% implied by the second specification. There is a similar analysis for innumerable other specifications, like (p = 0.04, $\hat{\theta} = 0.20$) and (p = 0.11, $\hat{\theta} = 0.26$) for example, where the posterior probabilities of an effect that is close to zero are respectively 15.1% and 14.9%. There are innumerable other such comparisons, with the footnote giving a few more examples.¹² In these comparisons, the specification with the larger *p*-value has a smaller (or equal) probability of the treatment effect being close to zero. Of course, in other comparisons, the specification with the larger *p*-value has a larger probability of the treatment effect being close to zero.

This illustrates why the results suggest caution about the practice of using small p-values as a key standard for empirical research. A larger (respectively, smaller) p-value does not necessarily correspond to a higher (respectively, lower) probability that θ is close to c. Comparing two studies, a study reporting a larger p-value can actually have a smaller posterior probability of a treatment effect that is close to 0. In particular, this shows that the research community can "miss" (e.g., not publish) treatment effects that are probably *not* close to zero if the research community only "accepts" (or focuses on) treatment effects with small p-values for the hypothesis of zero treatment effect. In particular, contrary to common usage, a larger p-value ("not rejecting") does not mean that there is a lack of evidence against the hypothesis that the treatment effect is (close to) zero. Rather than evaluate research based on the p-value, it is possible to use the expressions derived in this paper to assess the

¹²Consider these comparisons where the first specification has the smaller *p*-value, which implies rejecting the hypothesis that the effect is zero by classical standards and a smaller minimum posterior probability that the effect is zero. Compare $(p = 0.04, \hat{\theta} = 0.16)$ and $(p = 0.11, \hat{\theta} = 0.24)$: the respective posterior probabilities that the treatment effect is close to zero are 22.0% and 16.4%. Compare $(p = 0.04, \hat{\theta} = 0.12)$ and $(p = 0.90, \hat{\theta} = 0.12)$: the respective posterior probabilities that the treatment effect is close to zero are 36.6% and 8.3%. Compare $(p = 0.0001, \hat{\theta} = 0.08)$ and $(p = 0.11, \hat{\theta} = 0.12)$: the respective posterior probabilities that the treatment effect is close to zero are 83.5% and 39.3%.

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posterior probability that θ is close to c, which indeed is precisely what was done in this section. These remarks are general to empirical research, as a general feature of p-values, and are not particularly tied to the features of any specific empirical question or literature.

Remark 3 (Confidence intervals and standard errors). Some journals are now either recommending or requiring the use of alternative inferential statistics. For instance, the journals published by the American Economic Association (i.e., the American Economic Review, the American Economic Review: Insights, and the four American Economic Journals) and Econometrica and Quantitative Economics all ask authors to report standard errors, and not use asterisks to indicate statistical significance. Econometrica and Quantitative Economics further asks authors to report confidence intervals or coverage sets.¹³ Despite these prominent examples, which may represent the beginning of a general shifting of standards, currently journal policies against the use of p-values or statistical significance appear to be relatively rare overall in the scientific literature, per Hardwicke, Salholz-Hillel, Malički, Szűcs, Bendixen, and Ioannidis (2023).

Shifting from reporting p-values to reporting standard errors and/or confidence intervals does not avoid the problem with p-values. Reporting a p-value is "equivalent" to reporting a confidence interval, which in turn is "equivalent" to reporting a standard error, in the sense that any one such inferential quantity can be converted to the other inferential quantities, and therefore reporting any one of them conveys exactly the same information to the reader.¹⁴ Of course, it is not clear necessarily how *every* reader uses and interprets these other inferential quantities, but it seems clear that many readers use them in a way that directly parallels

 $^{^{13}\}mathrm{All}$ of this is clear from the online submission guidelines.

¹⁴This is because of the ability to convert between *p*-values for $\theta = c$ and confidence intervals for θ and standard errors for $\hat{\theta}_N$, for given $\hat{\theta}_N$. A $1-\alpha$ confidence interval has the form $[\hat{\theta}_N - c.v._{\alpha} \times s.e.(\hat{\theta}_N), \hat{\theta}_N + c.v._{\alpha} \times s.e.(\hat{\theta}_N)]$, where c.v._{α} is the appropriate critical value. This relationship allows a reader to convert between the confidence interval and the standard error, for given $\hat{\theta}_N$, so reporting a confidence interval conveys exactly the same information as does reporting a standard error. Further, the standard formula for the *p*-value gives a relationship that allows a reader to convert between the *p*-value and the standard error, for given $\hat{\theta}_N$, so reporting a *p*-value conveys exactly the same information as does reporting a standard error.

the use of p-values discussed throughout this paper, and therefore shares the same issues concerning the use of p-value that have been the focus of this paper.

It seems there are two ways that typical readers tend to use standard errors. First, a reader might think about the relative magnitude of $\hat{\theta}_N$ compared to the standard error. Specifically, a reader might think about whether the relative magnitude exceeds 1.96, which of course is equivalent to checking whether the *p*-value is less than 0.05. And, therefore, this use of standard errors results in the same issues concerning the use of *p*-values that have been the focus of this paper.

Alternatively, a reader can use a standard error to compute a confidence interval. For instance, a reader might think about (roughly what happens when) adding/subtracting a multiple (e.g., 1.96) of the standard error to/from $\hat{\theta}_N$. Then, a reader might think about whether c is contained within the confidence interval. This use of standard errors also is effectively the same as checking whether the p-value is less than the significance level that corresponds to the level of the confidence interval, which again results in the same issues concerning the use of p-values that have been the focus of this paper.

More generally, the results and discussions also apply to the relationship between confidence intervals and posterior probabilities of ϵ -approximate hypotheses. To illustrate the same issues arise, consider these specifications of 95% confidence intervals for a treatment effect: [0.0657, 0.1704], [0.0324, 0.2386], [0.0180, 0.2681], [-0.0193, 0.3425], [-0.0648, 0.4230]. Clearly, these are increasingly large (and nested) 95% confidence intervals. In order, the associated *p*-values for the hypothesis of zero treatment effect range from 0.00001 to 0.15. By the application of common empirical standards, essentially the same used above for *p*-values, the first specification easily meets the standard for statistical significance and rejecting the hypothesis that the treatment effect is zero, whereas the last specification does not achieve statistical significance and so it might be comparatively difficult to "sell"/publish as evidence of an effect. Because of the failure to reject the hypothesis of zero treatment effect, it might be said that the last specification shows "lack of evidence" against the hypothesis that the treatment effect is zero. What would be the corresponding posterior probabilities that the treatment effect is close to zero, which as before is defined for this example to be an effect in [-0.1, 0.1]? Actually, the posterior probabilities that the treatment effect is close to zero are exactly the same for these specifications: 25%. This reflects exactly the same issues discussed previously about *p*-values, because of the relationship between *p*-values and confidence intervals.

Remark 4 (Other approaches). This remark is about other approaches. Another case is when there is positive prior probability of $\theta = c$. As discussed in the Introduction, this case seems rare in empirical work in the social sciences. Nevertheless, this case can be analyzed. In this case, the posterior probability of $\theta \in [c - \epsilon, c + \epsilon]$ with ϵ small is approximately the same as the posterior probability of the hypothesis $\theta = c$ (e.g., Berger and Sellke (1987, Section 4)). Therefore, results for the minimum posterior probability of the hypothesis $\theta = c$ and of the hypothesis $\theta \in [c - \epsilon, c + \epsilon]$ with ϵ small are basically the same in this setup. In fact, this is part of the motivation for the literature on minimum posterior probabilities to study the (minimum) posterior probability of $\theta = c$, understanding it can be an approximation to the (minimum) posterior probability that $\theta \in [c - \epsilon, c + \epsilon]$. See for example the discussion in Berger and Sellke (1987). As mentioned in the Introduction, this means that the use of small *p*-values as a key standard, including *p*-value cutoff rules for significance like the one suggested by Benjamin, Berger, Johannesson, Nosek, Wagenmakers, et al. (2018), can be justified from a Bayesian perspective even if the empirical question is whether θ is close to *c*, when there is positive prior probability that $\theta = c$.

This paper studies the *p*-value for the hypothesis $\theta = c$, precisely because that is what is used in empirical practice, but one could in principle conduct a classical hypothesis test of the ϵ -approximate hypothesis (when ϵ is not data-dependent). In the appendix, Abadie (2020) studies the informativeness of rejecting or not rejecting the hypothesis $\theta \in [c - \epsilon, c + \epsilon]$ by such a classical hypothesis test. In large samples, Abadie (2020)'s results imply the posterior probability of the ϵ -approximate hypothesis is 1 conditional on not rejecting and is 0 conditional on rejecting.

Or, it is possible to conduct a classical hypothesis test of the one-sided hypothesis that $\theta \geq c$. In contrast to the motivating results cited in the introduction that concern the "calibration" of a *p*-value via the minimum posterior probability when the hypothesis is $\theta = 0$, when the hypothesis is $\theta \geq 0$, Casella and Berger (1987) show that the *p*-value for $\theta \geq c$ can exactly equal the minimum posterior probability that $\theta \geq c$ for certain classes of priors. Also, in an asymptotic setup similar to that used in this paper, Kline (2011) shows the same result.

The emphasis on the posterior probability of the ϵ -approximate hypothesis follows the related literature that also focuses on the posterior probability of certain hypotheses. However, it is possible to use these results to analyze other quantities; for instance, the posterior odds ratio is simply a (monotone) function of the posterior probability of the hypothesis.

Remark 5 (Data-dependent ϵ). ϵ can be data-dependent. For example, in a regression setting (with c = 0), ϵ can be a multiple λ of the estimated coefficient $\hat{\beta}$ on a "reference" explanatory variable. The ϵ -approximate hypothesis would be $\theta \in [-\lambda|\hat{\beta}|, \lambda|\hat{\beta}|]$, meaning the effect of one explanatory variable is "close to 0" if it is sufficiently closer to 0 compared to the effect of the "reference" explanatory variable. More specifically, consider as an example the case that θ reflects the treatment effect of a novel treatment, when the outcome is known already in the literature to be impacted by other "reference" treatments. Consider a regression model that includes both the novel treatment and one such "reference" treatment. ϵ can be selected to be proportionate to the estimated treatment effect of the reference treatment, so the ϵ -approximate hypothesis concerns whether or not the novel treatment has an effect that is at least a pre-specified percentage of the effect of the reference treatment. If so, the novel treatment can be said to have an "economically significant" effect, specifically relative to the effect of this reference treatment. Other choices of ϵ are allowed, including fixed constants as in the numerical example above. Another illustration of the choice of ϵ is provided in the empirical application in Section 5.

Remark 6 (Repeated sampling behavior). Another perspective comes from inspecting the repeated sampling behavior of the posterior probability of the ϵ -approximate hypothesis. The (large sample approximation to the) posterior probability of the ϵ -approximate hypothesis is a function of $\hat{\theta}_N$ and the *p*-value. Based on a given data generating process, it is possible to generate a sampling distribution of draws of $\hat{\theta}_N$ and the *p*-value, and for each such draw, compute the associated posterior probability of the ϵ -approximate hypothesis, thereby generating a sampling distribution of the posterior probability of the ϵ -approximate hypothesis.

Consider specifically the case that $X_i \sim^{iid} \mathcal{N}(\theta_0, \sigma_0^2)$, with $\hat{\theta}_N$ the associated sample average, based on N = 500. This data generating process is considered in the finite sample results in Section 6. The posterior probability of the ϵ -approximate hypothesis reported here comes from the finite sample results, but as discussed in Section 6, the results would be nearly identical numerically if the asymptotic approximation to the posterior probability were to be used. As in the above discussion, $\epsilon = 0.1$ for these calculations. The results are displayed in Figure 3.

Suppose first that $\theta_0 = 0$ and $\sigma_0^2 = 1$. Among the draws from the sampling distribution that "achieve statistical significance" with a *p*-value of 0.05 or below, considering the associated (conditional) sampling distribution of the posterior probability of the ϵ -approximate hypothesis, the 1st quantile is 0.11 and the 99th quantile is 0.64. There is a range both because a range of *p*-values is considered and because the posterior probability depends on $\hat{\theta}_N$, even for a given *p*-value. The quantiles of the sampling distribution are reported, rather than the support of the sampling distribution, because the sampling distribution has a particularly long tail of "unlikely" but extreme values of the posterior probabilities, a consequence of the expression for the posterior probability and the support of $\hat{\theta}_N$ and the *p*-value.



FIGURE 3. Repeated sampling density of the posterior probability of the ϵ -approximate hypothesis that $\theta \in [-0.1, 0.1]$.

Alternatively, suppose that $\theta_0 = 0$ and $\sigma_0^2 = 10$. For these purposes, an increase in σ_0^2 is similar to a decrease in N. Now, among the draws from the sampling distribution that "achieve statistical significance" with a p-value of 0.05 or below, considering the associated (conditional) sampling distribution of the posterior probability of the ϵ -approximate hypothesis, the 1st quantile is essentially 0 and the 99th quantile is 0.10.

This further illustrates, now in this repeated sampling sense, that caution is warranted about using "statistical significance" as a key standard in empirical research. Even just from these two data generating processes, out of the universe of data generating processes that could have generated the data, "statistical significance" is associated (in this repeated sampling sense) with a substantial range of associated posterior probabilities of the ϵ -approximate hypothesis.

Further, among the draws from the sampling distribution when $\sigma_0^2 = 1$ that do not "achieve statistical significance" with a *p*-value of above 0.05, considering the associated (conditional) sampling distribution of the posterior probability of the ϵ -approximate hypothesis, the 1st quantile is 0.64 and the 99th quantile is 0.98. And among the draws from the sampling distribution when $\sigma_0^2 = 10$ that do not "achieve statistical significance" with a *p*-value of above 0.05, considering the associated (conditional) sampling distribution of the posterior probability of the ϵ -approximate hypothesis, the 1st quantile is 0.11 and the 99th quantile is 0.54.

Thus, as also evident from Figure 3, the sampling distribution of the posterior probability of the ϵ -approximate hypothesis "without statistical significance" when $\sigma_0^2 = 10$ tends actually to have somewhat smaller values, compared to the sampling distribution of the posterior probability of the ϵ -approximate hypothesis "with statistical significance" when $\sigma_0^2 = 1$. This further illustrates that caution is warranted about using "statistical significance" as a key standard in empirical research. Although this discussion has focused on just two possible data generating processes, the basic idea would apply more broadly. Indeed, including additional data generating processes in the universe that are considered would further exacerbate the issues discussed above with just two data generating processes. For instance, this is apparent from panel 3b, which shows the case for $\theta_0 = 0.14$, where the sampling distribution of the posterior probability of the ϵ -approximate hypothesis tends to have smaller values compared to what happens when $\theta_0 = 0$, for any given combination of σ_0^2 and "statistical significance" (or not).

5. Empirical application

To further illustrate the results, this section presents an application of the results to published research from psychology in Open Science Collaboration (2015) and economics in Camerer, Dreber, Forsell, Ho, Huber, Johannesson, Kirchler, Almenberg, Altmejd, Chan, Heikensten, Holzmeister, Imai, Isaksson, Nave, Pfeiffer, Razen, and Wu (2016). The main goal of this empirical application is to show that there is a wide "range" of posterior probabilities of the ϵ -approximate hypothesis associated with any given *p*-value in actual empirical research, thereby reinforcing the caution against using *p*-values as a key standard for empirical research.

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Both of these investigations concern the "reproducibility" of published research. The original studies are selected because of their prominence, particularly in terms of the prominence of the journal. Each study is represented by one "key result" or "key effect" and a corresponding p-value, determined as part of the replication analysis. The estimated effects are correlation coefficients. Consequently, all of the estimated effects are mutually comparable. For effect sizes measured by correlation coefficients, the hugely influential Cohen (1969)¹⁵ has a lengthy discussion that states that an effect size of 0.1 is "small," an effect size of 0.3 is "medium," and an effect size of 0.5 is "large."

Summarized to the core feature that is relevant for the purposes of this paper, both investigations have two sets of estimated effects and p-values.¹⁶ One set comes from the original studies, and the other set comes from the corresponding replication studies. Each original study provides one estimated effect and one p-value; each replication study does the same. Given the inclusion criteria for these reproducibility investigations, this set of p-values is representative of "important" research.

Figure 4 shows the empirical scatter plot of the *p*-values and the corresponding posterior probability of the ϵ -approximate hypothesis, where ϵ can be 0.1 ("small") or 0.3 ("medium") or 0.5 ("large"). The results are displayed separately for the replication studies and the original studies, given the possible concerns with the original studies raised by the replication analysis. The much larger number of psychology studies are displayed in red circles, and the smaller number of economics studies are displayed in blue squares. Each mark corresponds to

¹⁵This has been cited approximately 240,000 times according to Google Scholar at the time of this writing. ¹⁶These investigations report the nature of the underlying hypothesis test, and those that evidently do not fit the framework of this paper are dropped. In practice, this primarily entails dropping *p*-values based on an *F*-test with strictly more than 1 numerator degrees of freedom. This arises particularly in the psychology investigation, from the application of two-way or multi-way ANOVA, which essentially amounts to (effects and) hypotheses involving multiple parameters of interest, rather than a single scalar parameter of interest as covered by this paper. (Of course, the analysis in this paper allows the underlying statistical model to have multiple parameters, but the analysis in this paper is for a scalar parameter of interest.) Given the nature of the analysis conducted here, which asks about the "range" of posterior probabilities of the ϵ -approximate hypothesis, this restriction only serves to bias "against" the finding that empirically there is a substantial range of posterior probabilities associated with any given *p*-value.

a study. In every panel of Figure 4, there is a substantial range of posterior probabilities of the ϵ -approximate hypothesis associated with any given *p*-value. Thus, among different studies with a similar *p*-value, there are very different posterior probabilities of the ϵ -approximate hypothesis. This further reinforces caution against using *p*-values as a key standard for empirical research. There are many pairs of studies displayed in Figure 4 such that the study with a higher *p*-value actually has a lower posterior probability of the ϵ -approximate hypothesis.

6. Finite sample results

A similar analysis can be done in finite samples, with additional distributional assumptions on the data generating process. $t_d(\mu, \Sigma)$ is a t-distribution with location μ , scale Σ , and degrees of freedom d. Thus $t_d(0, 1) \equiv t_d$ is the standard t-distribution with d degrees of freedom. \mathcal{F}_{d_1,d_2} is an F-distribution with degrees of freedom d_1 and d_2 . Define $G_a(v,w) \equiv$ $F_{t_a}\left(\sqrt{Q_{\mathcal{F}_{1,a}}(1-v)}(-1+w)\right) - F_{t_a}\left(\sqrt{Q_{\mathcal{F}_{1,a}}(1-v)}(-1-w)\right)$ for $a \in \mathbb{N}$ and $v \in (0,1)$. As with, essentially, any finite sample inference result (frequentist or Bayesian), the results in this section rely on a distributional assumption.

Assumption 2. It holds that:

- (1) The classical sampling distribution of $\frac{\hat{\theta}_N \theta_0}{\hat{\sigma}_N}$ has distribution $t_{N-d}(0, 1)$, i.e., a standard t-distribution with N-d degrees of freedom, for some estimated scale factor $\hat{\sigma}_N > 0$ and integer d such that N-d > 0.
- (2) The Bayesian posterior $\theta | X^{(N)}$ has distribution $t_{N-d}(\hat{\theta}_N, \hat{\sigma}_N^2)$.

An example of Assumption 2 is when $X_i \sim^{i.i.d.} \mathcal{N}(\theta, \sigma^2)$ with θ and σ^2 unknown. Then, $d = 1, \ \hat{\theta}_N = \frac{1}{N} \sum_{i=1}^N X_i$, and $\hat{\sigma}_N^2 = \frac{1}{N} \left(\frac{1}{N-1} \sum_{i=1}^N (X_i - \hat{\theta}_N)^2 \right)$. The prior is the typical improper "uninformative" uniform prior on $(\theta, \log \sigma)$. See Gelman, Carlin, Stern, Dunson, Vehtari, and Rubin (2013, Section 3.2). Another example is when θ is a linear regression coefficient with homoskedastic, normally-distributed unobservables. Then, d is the number



FIGURE 4. Empirical scatter plot of the posterior probability of the ϵ -approximate hypothesis that $\theta \in [-\epsilon, \epsilon]$ and the *p*-value.

of explanatory variables (counting the intercept), $\hat{\theta}_N$ is the OLS estimate, and $\hat{\sigma}_N^2$ is the estimate of the variance of the OLS estimate. The prior is a typical improper "uninformative" uniform prior. See Gelman, Carlin, Stern, Dunson, Vehtari, and Rubin (2013, Section 14.2) or Lancaster (2004, Section 3.3.3). As with the large sample approximation, and the discussion of Assumption 1, note again the analysis concerns the case of unknown variance (or scale) with corresponding estimate $\hat{\sigma}_N^2$. This assumption parallels typical assumptions used for classical inference in finite samples.

The classical test statistic is $W_N = \left(\frac{\hat{\theta}_N - c}{\hat{\sigma}_N}\right)^2$. By Assumption 2, if the hypothesis $\theta = c$ is true, W_N has distribution $\mathcal{F}_{1,N-d}$. Therefore, the *p*-value is $p_N = \overline{F}_{\mathcal{F}_{1,N-d}}(W_N)$.

Theorem 2. Under Assumption 2, and when the p-value is not $1,^{17}$ for any $\epsilon > 0$ chosen by the econometrician (possibly data-dependent), the Bayesian posterior probability of the ϵ -approximate hypothesis that $\theta \in [c - \epsilon, c + \epsilon]$ is $G_{N-d}\left(p_N, \frac{\epsilon}{|\hat{\theta}_N - c|}\right)$.

Lemma 1. $G_a(v,w) \to F_{\chi^2_{1,Q_{\chi^2_1}(1-v)}}\left(w^2 Q_{\chi^2_1}(1-v)\right) as a \to \infty.$

As $N \to \infty$, Theorem 2 limits to Theorem 1 by applying Lemma 1. In fact, Theorem 2 and Theorem 1 are practically the same numerically when $N - d \ge 1000$, and also when $N - d \ge 300$ and $p_N \ge 0.001$, among other cases.¹⁸

This shows that the main point of this paper is not tied to a particular setup (e.g., "asymptotic approximation" or "finite sample") but rather a general feature of *p*-values. This also reinforces that the large sample approximation is achieving a valid approximation to finite sample behavior, based on the non-degenerate approximation to the posterior distribution that is discussed after the statement of Theorem 1.

¹⁷Similar to the asymptotic approximation, $p_N = 1$ happens with probability 0 when $\hat{\theta}_N$ has a continuous sampling distribution and $p_N = 0$ cannot happen in this setup.

¹⁸This claim is based on the maximal absolute difference between the posterior probability of the ϵ -approximate hypothesis per Theorem 2 and the asymptotic approximation, where the maximum is across values of p_N , ϵ , and $\hat{\theta}_N$. In the stated cases, the maximal absolute difference is less than 0.003.

Figure 5 shows the posterior probability of the ϵ -approximate hypothesis, for varying values of N-d. Except for extremely small values of N-d, the result is almost identical to the large sample approximation in Figure 1, which appears also as Figure 5f. The main qualitative features of Figure 1 (or equivalently Figure 5f) are also present in the finite sample panels of Figure 5.

7. Conclusions

This paper derives closed-form expressions for the posterior probability of the ϵ -approximate hypothesis. The properties are discussed, which contradict some common practices about using *p*-values. In the running example of treatment effects, an implication of the results is that larger *p*-values can correspond to lower posterior probabilities that the treatment effect is close to zero, potentially lower than from a smaller *p*-value. This shows that the research community can "miss" (e.g., not publish) treatment effects that are probably *not* close to zero if the research community only "accepts" (or focuses on) treatment effects with small *p*-values for the hypothesis of zero treatment effect. In particular, contrary to common usage, a larger *p*-value ("not rejecting") does not mean that there is a lack of evidence against the hypothesis that the treatment effect is (close to) zero. On the other hand, it is possible to use the expressions derived in this paper to assess the posterior probability of ϵ -approximate hypotheses even without access to the original data, in the sense that the expressions depend only on quantities that are conventionally reported in research, thereby assessing the evidence about whether the treatment effect is close to zero.

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FIGURE 5. Posterior probability of the ϵ -approximate hypothesis that $\theta \in [c - \epsilon, c + \epsilon]$ for given *p*-value and $\frac{|\hat{\theta}_N - c|}{\epsilon}$ according to the finite sample result with varying N - d, and large sample approximation.

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Appendix A. Technical appendix

Lemma 2. Under Assumption 1, $||\Pi_{\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} - \mathcal{N}(0,\hat{\Sigma}_N)||_{TV} \rightarrow^p 0$ and $||\Pi_{\hat{\Sigma}_N^{-\frac{1}{2}}\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} - \mathcal{N}(0,I_{m\times m})||_{TV} \rightarrow^p 0.$

Proof of Lemma 2. $||\mathcal{N}(0,\hat{\Sigma}_N) - \mathcal{N}(0,\Sigma_0)||_{TV} \to^p 0$ since total variation distance is bounded above by the square root of Kullback-Leibler divergence (e.g., DasGupta (2008, Chapter 2)) and $\hat{\Sigma}_N \to^p \Sigma_0$ by Assumption 1. Therefore, $||\Pi_{\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} - \mathcal{N}(0,\hat{\Sigma}_N)||_{TV} \to^p 0$.

Let $h_N(\cdot) : \mathbb{R}^m \to \mathbb{R}^m$ be $h_N(w) = \hat{\Sigma}_N^{-\frac{1}{2}} w$. Since $h_N(\cdot)$ is continuous, $h_N^{-1}(\mathcal{B})$ is Borel for any Borel \mathcal{B} . Therefore, $|\Pi_{\hat{\Sigma}_N^{-\frac{1}{2}}\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}}(\mathcal{B}) - \mathcal{N}(0, I_{m \times m})(\mathcal{B})| = |\Pi_{\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}}(h_N^{-1}(\mathcal{B})) - \mathcal{N}(0, \hat{\Sigma}_N)(h_N^{-1}(\mathcal{B}))|$. Because $||\Pi_{\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} - \mathcal{N}(0, \hat{\Sigma}_N)||_{TV} \to^p 0$, $||\Pi_{\hat{\Sigma}_N^{-\frac{1}{2}}\sqrt{N}(\psi-\hat{\psi}_N)|X^{(N)}} - \mathcal{N}(0, I_{m \times m})||_{TV} \to^p 0$.

For the hypothesis $\tilde{\psi} = c$, where $\tilde{\psi} \in \mathbb{R}^{\tilde{m}}$ is a sub-vector of components of ψ (e.g., $\tilde{\psi} = \theta$ or $\tilde{\psi} = (\theta, \gamma_2)$), and $\tilde{\Sigma}$ is the corresponding sub-matrix of Σ , the test statistic is

$$\tilde{W}_N = N(\hat{\tilde{\psi}}_N - c)'\hat{\tilde{\Sigma}}_N^{-1}(\hat{\tilde{\psi}}_N - c).$$

The p-value is

$$\tilde{p}_N = P(\chi^2_{\tilde{m}} > \tilde{W}_N) = \overline{F}_{\chi^2_{\tilde{m}}}(\tilde{W}_N).$$

Lemma 3. Under Assumption 1,

$$\sup_{0 < q < 1} \left| q - \Pi \left(\delta_{\hat{\Sigma}_{N,2}}(\tilde{\psi}, c) \le \sqrt{\frac{Q_{\chi^2_{\tilde{m}, Q_{\chi^2_{\tilde{m}}}(1 - \tilde{p}_N)}(q)}}{N} \mid X^{(N)}} \right) \right| \to^p 0.$$

Proof of Lemma 3.
$$||\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)||_{2}^{2} = \sqrt{N}(\tilde{\psi}-c)'\hat{\Sigma}_{N}^{-1}\sqrt{N}(\tilde{\psi}-c) = \sqrt{N}(\tilde{\psi}-\hat{\psi}_{N}+\hat{\psi}_{N}+\hat{\psi}_{N}+c)|_{2}^{2}$$
. Let $Z \sim \mathcal{N}(0, I_{\tilde{m}\times\tilde{m}})$. For any Borel $\mathcal{B} \subseteq \mathbb{R}^{\tilde{m}}$, $\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)|X^{(N)}}(\mathcal{B}) = \Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-\hat{\psi}_{N}+\hat{\psi}_{N}-c)|X^{(N)}}(\mathcal{B}) = \Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-\hat{\psi}_{N})|X^{(N)}}(\mathcal{B}-\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c)) - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))(\mathcal{B})| = |\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-\hat{\psi}_{N})|X^{(N)}}(\mathcal{B}-\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c)) - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))(\mathcal{B})| = |\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-\hat{\psi}_{N})|X^{(N)}}(\mathcal{B}-\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c)) - Z(\mathcal{B}-\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))| \rightarrow^{p} 0$. Therefore, $||\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)|X^{(N)}} - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))||_{TV} \rightarrow^{p} 0$. Let $h(\cdot) : \mathbb{R}^{\tilde{m}} \rightarrow \mathbb{R}$ be given by $h(z) = z'z$. Since $h(\cdot)$ is continuous, $h^{-1}(\mathcal{B}) \subseteq \mathbb{R}^{\tilde{m}}$ is Borel for any Borel \mathcal{B} . Therefore, $||\Pi_{||\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)||_{2}^{2}|X^{(N)}}(h^{-1}(\mathcal{B})) - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))(h^{-1}(\mathcal{B}))|$. Because $||\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)|X^{(N)}} - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))(h^{-1}(\mathcal{B}))|$. Because $||\Pi_{\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}_{N}-c)|X^{(N)}} - (Z+\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\hat{\psi}_{N}-c))||_{TV} \rightarrow^{p} 0$.

$$C_N = (Z + \hat{\Sigma}_N^{-\frac{1}{2}} \sqrt{N}(\hat{\psi}_N - c))' (Z + \hat{\Sigma}_N^{-\frac{1}{2}} \sqrt{N}(\hat{\psi}_N - c)) \sim \chi^2_{\hat{m}, \tilde{W}_N}.$$
 Therefore,

 $\sup_{0 < q < 1} |\Pi_{||\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)||_{2}^{2}|X^{(N)}}((-\infty, Q_{\chi^{2}_{\tilde{m},Q_{\chi^{2}_{\tilde{m}}}(1-\tilde{p}_{N})}}(q)]) - C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},Q_{\chi^{2}_{\tilde{m}}}(1-\tilde{p}_{N})}}(q)])| \to^{p} 0.$ By definition, $\tilde{W}_{N} = Q_{\chi^{2}_{\tilde{m}}}(1-\tilde{p}_{N})$, so $C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},Q_{\chi^{2}_{\tilde{m}}}(1-\tilde{p}_{N})}}(q)]) = C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},\tilde{W}_{N}}}(q)]) = C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},\tilde{W}_{N}}}(q)]) = C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},\tilde{W}_{N}}}(q)]) = C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},\tilde{W}_{N}}}(q)]) = C_{N}((-\infty, Q_{\chi^{2}_{\tilde{m},\tilde{W}_{N}}}(q)))$

 $q. \text{ The result follows since by definition } \Pi(\delta_{\hat{\Sigma}_{N},2}^{2}(\tilde{\psi},c) \leq \frac{Q_{\chi_{\tilde{m},Q_{\chi_{\tilde{m}}^{2}(1-\tilde{p}_{N})}}(q)}{N} \mid X^{(N)}) = \Pi_{||\hat{\Sigma}_{N}^{-\frac{1}{2}}\sqrt{N}(\tilde{\psi}-c)||_{2}^{2}|X^{(N)}}((-\infty,Q_{\chi_{\tilde{m},Q_{\chi_{\tilde{m}}^{2}(1-\tilde{p}_{N})}}(q)]).$

Lemma 4. Under Assumption 1,

$$\sup_{0 < q < 1} \left| \left(q - \Pi \left(\left| \theta - c \right| \le \sqrt{\frac{Q_{\chi_{1,Q_{\chi_{1}}^{2}(1-p_{N})}}(q)}{Q_{\chi_{1}^{2}}(1-p_{N})}} |\hat{\theta}_{N} - c| \mid X^{(N)} \right) \right) 1[p_{N} < 1] \right| = o_{p}(1).$$

Proof of Lemma 4. $Q_{\chi_1^2}(1-p_N) = W_N = N(\hat{\Sigma}_{N,11})^{-1}(\hat{\theta}_N - c)^2$, so $(\hat{\Sigma}_{N,11})^{-\frac{1}{2}} = \frac{\sqrt{Q_{\chi_1^2}(1-p_N)}}{\sqrt{N}|\hat{\theta}_N - c|}$ when $|\hat{\theta}_N - c| \neq 0$ which is equivalent to $p_N < 1$. When $|\hat{\theta}_N - c| \neq 0$, $\delta_{\hat{\Sigma}_{N,11},2}(\theta, c) = (\hat{\Sigma}_{N,11})^{-\frac{1}{2}}|\theta - c| = \frac{\sqrt{Q_{\chi_1^2}(1-p_N)}}{\sqrt{N}|\hat{\theta}_N - c|}|\theta - c|$. The result follows from Lemma 3 by substitution. \Box

$$\begin{array}{l} Proof \ of \ Theorem \ 1. \ \text{By simplification}, \ \Pi\left(\theta \in [c-\epsilon,c+\epsilon]|X^{(N)}\right) = \\ \Pi\left(|\theta-c| \leq \sqrt{\frac{Q_{\chi^2_{1,Q_{\chi^2_1}(1-p_N)}}\left(F_{\chi^2_{1,Q_{\chi^2_1}(1-p_N)}}\left(\frac{\epsilon^2}{|\theta_N-c|^2}Q_{\chi^2_1}(1-p_N)\right)\right)}{Q_{\chi^2_1}(1-p_N)}}|\hat{\theta}_N-c||X^{(N)}\right). \ \text{ Therefore, the result follows from Lemma 4.} \\ \end{array}$$

Proof of Corollary 1 and Corollary 2. By Johnson, Kotz, and Balakrishnan (1995, page 441), $F_{\chi_{1,b}^2}(ab)$ where $a = \frac{e^2}{|\theta_N - c|^2}$ and $b = Q_{\chi_1^2}(1 - p_N)$ can be written $\Phi(\sqrt{b}(\sqrt{a} - 1)) - \Phi(-\sqrt{b}(\sqrt{a} + 1))$. The derivative with respect to b is $\frac{1}{\sqrt{2\pi}}\exp(-\frac{1}{2}(ab - 2\sqrt{a}b + b))(\sqrt{a} - 1)\frac{1}{2}b^{-\frac{1}{2}} + \frac{1}{\sqrt{2\pi}}\exp(-\frac{1}{2}(ab + 2\sqrt{a}b + b))(\sqrt{a} + 1)\frac{1}{2}b^{-\frac{1}{2}}$. The sign of the derivative is the same as the sign of $\exp(2\sqrt{a}b)(\sqrt{a} - 1) + (\sqrt{a} + 1)$. If $a \ge 1$, the derivative is obviously positive. The first part of Corollary 2 follows, since b is a decreasing function of p_N . If a < 1, the derivative is negative (resp. positive, zero) exactly when b is greater than (resp. less than, equal to) $\frac{1}{2\sqrt{a}}\log\left(\frac{-\sqrt{a}-1}{\sqrt{a}-1}\right)$. The first part of Corollary 1 follows, since this means the posterior probability is a decreasing (resp., increasing) function of p_N when p_N is greater than (resp., less than) $1 - F_{\chi_1^2}\left(\frac{1}{2\sqrt{a}}\log\left(\frac{-\sqrt{a}-1}{\sqrt{a}-1}\right)\right)$. The limits in these Corollaries are apparent from the representation from Johnson, Kotz, and Balakrishnan (1995, page 441) that implies that $F_{\chi_{1,\ell^2}^2}(s^2t^2) = \Phi(t(s-1)) - \Phi(t(-s-1))$.

Proof of Theorem 2. Follow the strategy of Theorem 1. $W_N = Q_{\mathcal{F}_{1,N-d}}(1-p_N)$ and $\hat{\sigma}_N^{-1} = \frac{\sqrt{Q_{\mathcal{F}_{1,N-d}}(1-p_N)}}{|\hat{\theta}_N-c|}$. The posterior for $\hat{\sigma}_N^{-1}(\theta-c)$ is $t_{N-d}(\hat{\sigma}_N^{-1}(\hat{\theta}_N-c),1)$. Therefore, $\Pi(|\theta-c| \leq \epsilon |X^{(N)}) = \Pi(-\hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) \leq \hat{\sigma}_N^{-1}(\theta-c) - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) \leq \hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) \leq c_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) \leq t_{N-d} \leq \hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) = P(-\hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c) \leq t_{N-d} \leq \hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}(\hat{\theta}_N-c)) = P(-\hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}|\hat{\theta}_N-c| \leq t_{N-d} \leq \hat{\sigma}_N^{-1}\epsilon - \hat{\sigma}_N^{-1}|\hat{\theta}_N-c|)$ by symmetry of the t_{N-d} distribution. The final expression is $G_{N-d}(p_N, \frac{\epsilon}{|\hat{\theta}_N-c|})$ using the expression for $\hat{\sigma}_N^{-1}$.

Proof of Lemma 1. $\mathcal{F}_{1,a} \to \chi_1^2$ and $t_a \to \mathcal{N}(0,1)$ as $a \to \infty$. Then use the representation for $F_{\chi_{1,t^2}^2}(s^2t^2)$ used in the proof of Corollaries 1 and 2.

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