



SAPIENZA
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Bayesian Sample size for calibration of approximate intervals in clinical trials

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Outline

- ▶ Interval estimation of θ
- ▶ C : exact Highest Posterior Density (**HPD**) interval
- ▶ \tilde{C} : Likelihood Normal Approximate (**LNA**) interval

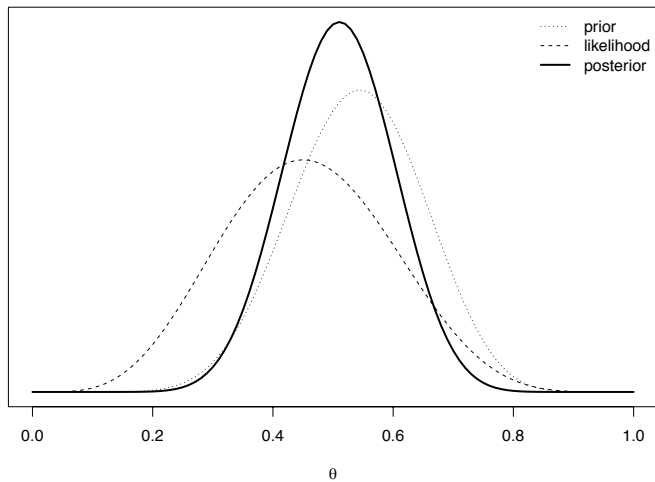
Problem: what n such that $\tilde{C} \simeq C$?

Model

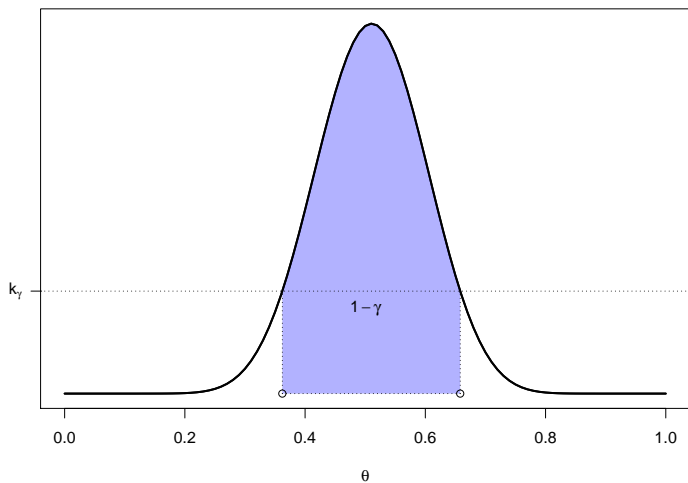
Beta-Binomial model

$$\left. \begin{array}{l} \mathbf{X}_n = (X_1, \dots, X_n) \\ X_i | \theta \sim \text{Ber}(\theta) \\ \theta \sim \text{Be}(\alpha, \beta) \end{array} \right\} \implies \theta | \mathbf{x}_n \sim \text{Be}(\bar{\alpha}, \bar{\beta})$$

Posterior distribution



Exact credible intervals for θ : HPD



Exact credible intervals for θ

$$\text{HPD: } C(\mathbf{x}_n) = [\ell(\mathbf{x}_n), u(\mathbf{x}_n)]$$

- ▶ $C(\mathbf{x}_n) = \{\theta \in \Theta : \pi(\theta|\mathbf{x}_n) \geq k_\gamma\},$
- ▶ $\mathbb{P}[\theta \in C(\mathbf{x}_n)|\mathbf{x}_n] = 1 - \gamma.$

Exact credible intervals for θ

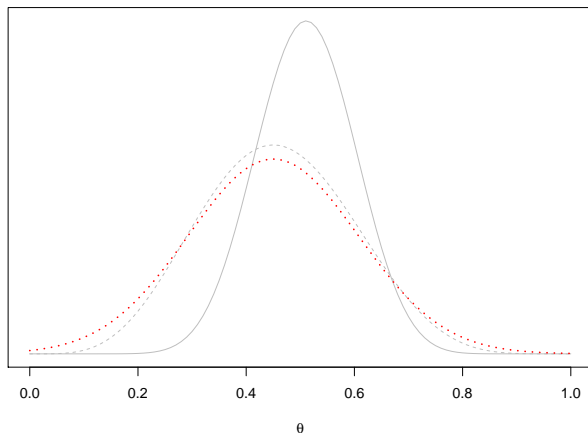
$$\text{HPD: } C(\mathbf{x}_n) = [\ell(\mathbf{x}_n), u(\mathbf{x}_n)]$$

- ▶ $(1 - \gamma)$ shortest intervals
- ▶ **but** no closed-form expression

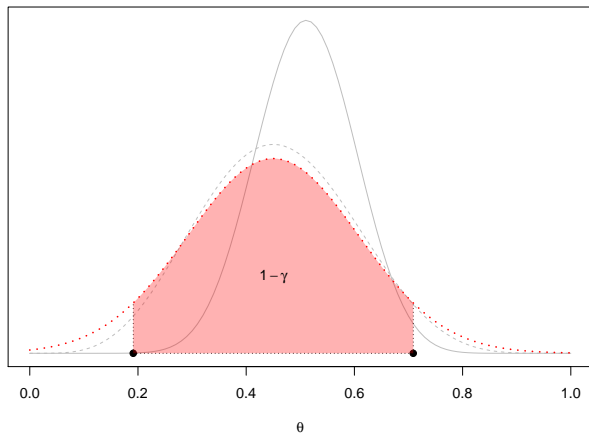
Likelihood Normal Approximation of the Posterior

$$\theta | \mathbf{x}_n \sim \mathcal{N}[\hat{\theta}, I_n(\hat{\theta})^{-1}]$$

with $\hat{\theta}$ MLE and $I_n(\theta)$ expected Fisher Information



Likelihood Normal Approximation of the Posterior



Approximate credible intervals for θ

$$\text{LNA: } \tilde{C}(\mathbf{x}_n) = [\tilde{\ell}(\mathbf{x}_n), \tilde{u}(\mathbf{x}_n)]$$

$$\blacktriangleright \tilde{\ell} = \hat{\theta} - z_{1-\frac{\gamma}{2}} I_n(\hat{\theta})^{-1/2} = \bar{x}_n - z_{1-\frac{\gamma}{2}} \sqrt{\bar{x}_n(1 - \bar{x}_n)/n}$$

$$\blacktriangleright \tilde{u} = \hat{\theta} + z_{1-\frac{\gamma}{2}} I_n(\hat{\theta})^{-1/2} = \bar{x}_n + z_{1-\frac{\gamma}{2}} \sqrt{\bar{x}_n(1 - \bar{x}_n)/n}$$

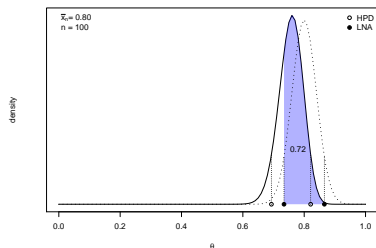
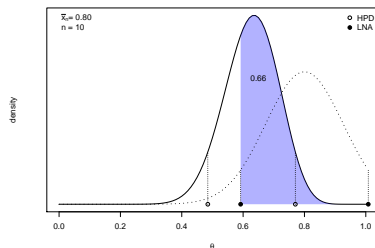
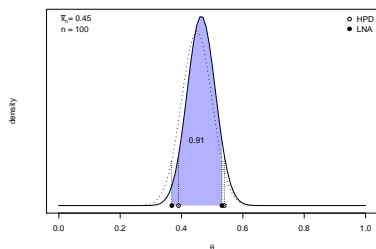
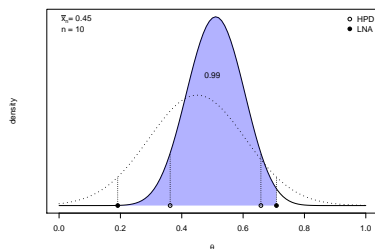
Calibrated approximate intervals

- ▶ In general, the *actual* posterior probability of \tilde{C} may be different from its *nominal* value $1 - \gamma$.
- ▶ We say \tilde{C} is *calibrated* if its exact posterior probability is equal to $1 - \gamma$.
- ▶ As n increases C and \tilde{C} converge, i.e.

$$\mathbb{P}[\theta \in \tilde{C} | \mathbf{x}_n] \rightarrow (1 - \gamma).$$

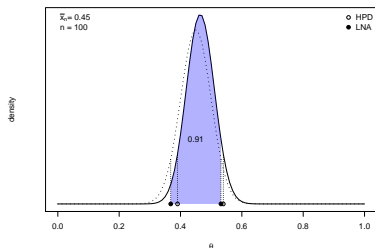
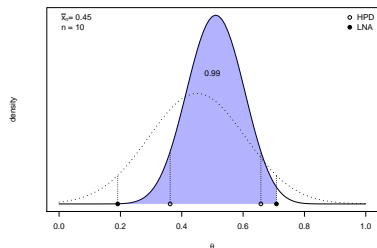
Posterior density VS its likelihood approximation

$1 - \gamma = 0.9$



Posterior density VS its likelihood approximation

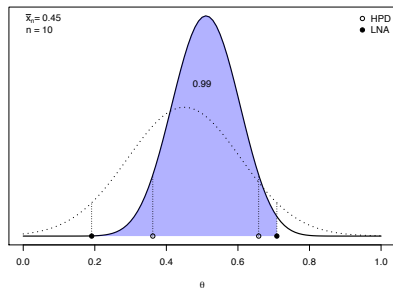
$$1 - \gamma = 0.9$$



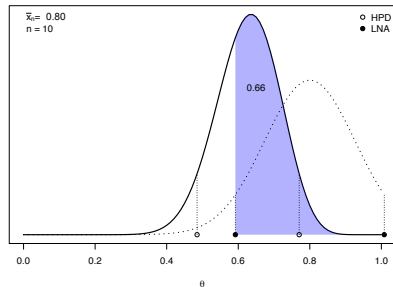
larger sample sizes ► better approximations

Posterior density VS its likelihood approximation

$$1 - \gamma = 0.9$$



larger distance between
likelihood mode (MLE)
and posterior mode



greater discrepancy between
HPD and LNA

Idea

A measure of lack of calibration of \tilde{C} , is:

$$|\mathbb{P}[\theta \in \tilde{C} | \mathbf{x}_n] - (1 - \gamma)|$$

- ▶ quantifies the discrepancy between the actual posterior probability of \tilde{C} and $(1 - \gamma)$
- ▶ ranges in $(0, 1 - \gamma)$

Hence, a **relative measure of discrepancy** is

$$P(\mathbf{x}_n) = \frac{|\mathbb{P}[\theta \in \tilde{C} | \mathbf{x}_n] - (1 - \gamma)|}{1 - \gamma}$$

Sample size determination

- ▶ Before observing the data, $P(\mathbf{X}_n)$ is random.
- ▶ The calibration of $\tilde{C}(\mathbf{X}_n)$ can be studied by looking at

$$e_n^P = \mathbb{E}_d[P(\mathbf{X}_n)]$$

computed w.r.t. the distribution of \mathbf{X}_n for a design value θ_d .

- ▶ To obtain a calibrated interval, select

$$n_P^* = \min\{n \in \mathbb{N} : e_n^P < \epsilon\}.$$

Computation of e_n^P

In simple cases with exact calculations...more often (*always*) via
Monte Carlo simulation

For each n , θ_d and $N = 10000$,

1. draw N samples $\mathbf{x}_n^{(1)}, \dots, \mathbf{x}_n^{(N)}$ from $f_n(\cdot; \theta_d)$;
2. compute $\tilde{\ell}(\mathbf{x}_n^{(j)})$ and $\tilde{u}(\mathbf{x}_n^{(j)})$, for $j = 1, \dots, N$;
3. compute $P(\mathbf{x}_n^{(j)})$, for $j = 1, \dots, N$;
4. set $e_n^P \simeq \frac{\sum_{j=1}^N P(\mathbf{x}_n^{(j)})}{N}$.

Example: single-arm phase II trial

- ▶ **Setup:** based on a trial conducted between 2009 and 2011 to test the combination of *lenalidomide and rituximab* in patients with *recurrent indolent non-follicular lymphoma* (see Sacchi et al, 2016, and Sambucini, 2019)
- ▶ **Goal:** interval estimation of the treatment response rate θ
- ▶ **Endpoint:** sample treatment response rate $\hat{\theta}$,
i.e. proportion of patients achieving complete/partial response

Example: single-arm phase II trial

- ▶ **Hystorical data:** 21 responses out of 39 patients
- ▶ **Prior elicitation:** $\theta \sim Be(\alpha, \beta)$ based on hystorical data
 \Rightarrow prior mean $\alpha/(\alpha + \beta) = 0.54$
- ▶ **Sensitivity analysis:** different amount of information
 \Rightarrow prior sample sizes $\alpha + \beta = (5, 10, 20)$
- ▶ **Non-informative case:** $\alpha = \beta = 1 \Rightarrow \theta \sim Unif(0, 1)$

Example: single-arm phase II trial

▶ **Design values:**

★ lowest acceptable value: $\theta_d = 0.45$

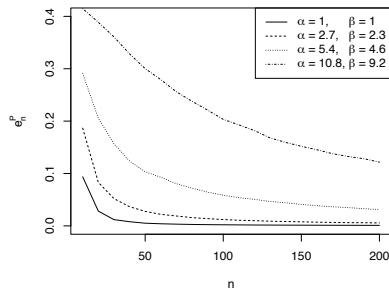
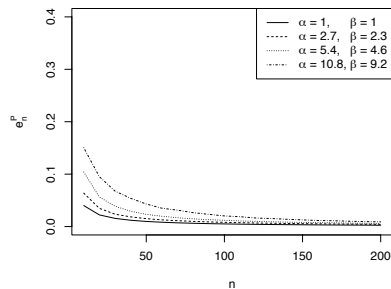
★ optimistic design value: $\theta_d = 0.80$

▶ **SSD criterion threshold:**

$$\epsilon = 0.01$$

Example: single-arm phase II trial

Plots of e_n^P w.r.t. n for several values of (α, β) with $\theta_d = 0.45$ and $\theta_d = 0.8$

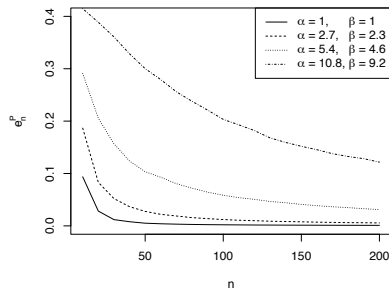
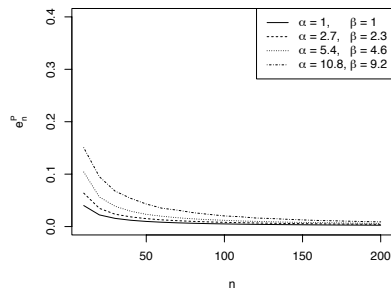


Remarks

1. *Effect of n .* Values of e_n^P decrease as n increases

Example: single-arm phase II trial

Plots of e_n^P w.r.t. n for several values of (α, β) with $\theta_d = 0.45$ and $\theta_d = 0.8$

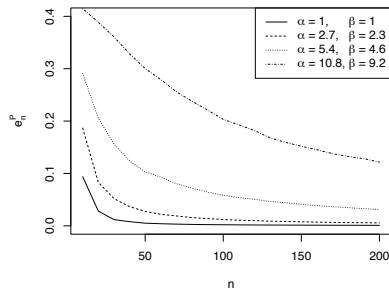
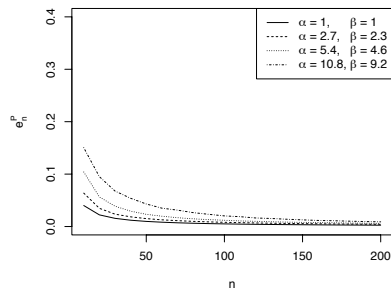


Remarks

2. *Effect of prior sample size.* Large $\alpha + \beta \Rightarrow$ large e_n^P

Example: single-arm phase II trial

Plots of e_n^P w.r.t. n for several values of (α, β) with $\theta_d = 0.45$ and $\theta_d = 0.8$



Remarks

3. *Effect of $|\theta_d - \mathbb{E}(\theta)|$.* If $\alpha + \beta \gg n$, large $|\theta_d - \mathbb{E}(\theta)| \Rightarrow$ large e_n^P

Example: single-arm phase II trial

Optimal sample sizes for different prior parameters, design values, $\epsilon_P = 0.01$, $\epsilon_L = 0.1$.

θ_d	(α, β)	(1, 1)	(2.7, 2.3)	(5.4, 4.6)	(10.8, 9.2)
0.45	n_P^*	49	80	119	182
	n_L^*	265	262	257	247
	\tilde{n}_L^*	267	267	267	267
0.80	n_P^*	35	118	646	2911
	n_L^*	170	169	169	167
	\tilde{n}_L^*	172	172	172	172

Example: single-arm phase II trial

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	n_L^*	170	169	169	167
	\tilde{n}_L^*	172	172	172	172

Remarks

4. *Comparison with ALC.* Opposite behaviour of n_P^* and n_L^* .
Large $\alpha + \beta \Rightarrow$ short intervals \Rightarrow small n_L^*

Comments · Objections · Extensions

The traditional way of using Bayesian intervals...

- ▶ pre-posterior control of length and position of credible interval

A different look at the problem...

- ▶ number of units needed to *safely* use LNA instead of HPD
- ▶ price of using LNA instead of HPD *in terms of expected discrepancy*

Our criterion can be put beside additional criteria related to the main goal of the trial

Comments · Objections · Extensions

One drawback of \tilde{C} is that it may include values outside $[0, 1]$

- ▶ Common solution: $\psi = g(\theta)$ (e.g. the logodds)
- ▶ Normal approximation improves
- ▶ Implementation is substantially similar...
 - ▷ HPD bounds are obtained via MC
 - ▷ Delta method provides closed form expressions of LNA
- ▶ Similar comments apply

References

Bayes for CT & SSD methods based on intervals

- ▶ Spiegelhalter et al (2004) [Bayesian Approaches to Clinical Trials and Health-Care Evaluation](#). Statistics in Practice. Wiley.
- ▶ Joseph L, Wolfson D B, Berger R Du (1995) [Sample Size Calculations for Binomial Proportions via Highest Posterior Density Intervals](#). J Roy Stat Soc: Series D (Statistician), 44: 143-154.
- ▶ M'Lan C E, Joseph L, Wolfson D B (2008) [Bayesian sample size determination for binomial proportions](#). Bayesian Anal, 3: 269-296.
- ▶ Cao J, Lee J Jack, Alber S (2009) [Comparison of Bayesian sample size criteria: ACC, ALC, and WOC](#). J Stat Plan Inference, 139: 4111-4122.
- ▶ Joseph L, Belisle P (2019) [Bayesian consensus-based sample size criteria for binomial proportions](#). Stat Med, doi: 10.1002/sim.8316.

References

Our contributions to interval-based Bayes SSD

- ▶ Brutti P, De Santis F (2008) Robust Bayesian sample size determination for avoiding the range of equivalence in clinical trials. J Stat Plan Inference, 138: 1577-1591.
- ▶ Gubbiotti S, De Santis F (2011) A Bayesian method for the choice of the sample size in equivalence trials. Aust N Z J Stat, 53: 443-460.
- ▶ Brutti P, De Santis F, Gubbiotti S (2014) Predictive measures of the conflict between frequentist and Bayesian estimators. J Stat Plan Inference, 148: 111-122.

References

Application

- ▷ Sacchi et al (2016) Safety and efficacy of lenalidomide in combination with rituximab in recurrent indolent non-follicular lymphoma: final results of a phase II study conducted by the Fondazione Italiana Linfomi *Haematologica*, 101 (5), e196-e199.
- ▷ Sambucini, V (2019) Bayesian predictive monitoring with bivariate binary outcomes in phase II clinical trials. *Computational Statistics & Data Analysis*, Elsevier, vol. 132(C), 18-30.