# Noninformative Priors for the Ratio of the Lognormal Means with Equal Variances

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#### Abstract

We develop noninformative priors for the ratio of the lognormal means in equal variances case. The Jeffreys' prior and reference priors are derived. We find a first order matching prior and a second order matching prior. It turns out that Jeffreys' prior and all of the reference priors are first order matching priors and in particular, one-at-a-time reference prior is a second order matching prior. One-at-a-time reference prior meets very well the target coverage probabilities. We consider the bioequivalence problem. We calculate the posterior probabilities of the hypotheses and Bayes factors under Jeffreys' prior, reference prior and matching prior using a real-life example.

Keywords: Bioequivalence problem; equal variance; Jeffreys' prior; matching priors; ratio of the lognormal means; reference priors.

#### 1. Introduction

Faced with a skewed distribution, one can transform the original data in a way that normalizes the distribution. The log-transformation is the most commonly used one. One estimation problem is to construct confidence intervals and to test for the lognormal mean. A common practice for such comparisons is to perform testing procedures on log-transformed outcome variables and to report the resulting p-values for the null hypothesis based on the original outcomes. For the Bayesian approach, see Moon  $et\ al.\ (2000)$  and Moon and Kim (2001). These proposed the methods which are based on the Bayes factor. Another estimation problem is for obtaining confidence intervals and tests for the ratio of means of two independent lognormal distributions. For example, in a bioequivalence

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trials, the relative potency of a new drug to that of a standard one is expressed in terms of the ratio of means, and analysts often need to construct a confidence interval for this ratio or to test the null hypothesis that the ratio is one (Berger and Hsu, 1996; Chow and Liu, 2000).

For our interest parameter, the ratio of lognormal means, we consider here noninformative priors. We consider Bayesian priors such that the resulting credible intervals for the ratio of the lognormal means have coverage probabilities equivalent to their frequentist counterparts. Although this matching can be justified only asymptotically, our simulation results indicate that this is indeed achieved for small or moderate sample sizes as well. The matching idea goes back to Welch and Peers (1963). Interest in such priors revived with the work of Stein (1985) and Tibshirani (1989). Among others, we may cite the work of Mukerjee and Dey (1993), Datta and Ghosh (1995a, 1995b), Datta and Ghosh (1995c, 1996), and Mukerjee and Ghosh (1997). On the other hand, Ghosh and Mukerjee (1992), and Berger and Bernardo (1989, 1992) extended Bernardo's (1979) reference prior approach, giving a general algorithm to derive a reference prior by splitting the parameters into several groups according to their order of inferential importance. This approach is very successful in various practical problems. Quite often reference priors satisfy the matching criterion described earlier.

In this paper, we develop noninformative priors when the variances of the logtransformed outcome variables are equal. In Section 2, we derive Fisher information matrix under orthogonal reparametrization. Then we develop the various noninformative priors. Specifically we derive the Jeffreys' prior and the reference priors for different groups of ordering for the parameters, and we develop first order and second order probability matching priors for our interesting parameter. Sufficient conditions for the propriety of posterior for a general class of priors are also given in this section. In Section 3, we provide the simulated frequentist coverage probabilities under the proposed priors. In Section 4, we consider the bioequivalence problem. Two different drugs or formulations of the same drugs are called bioequivalent if they are absorbed into the blood and become available at the drug action site at about the same rate and concentration. Bioequivalence of the two drugs is defined as the ratio of means is between the tolerance limit prespecified by a regulatory agency. We calculate the posterior probabilities of the hypotheses under Jeffreys' prior, reference prior and matching prior. Also we provide the Bayes factors. Two treatments AUC data are provided for equal variance case to illustrate our results.

## 2. Development of Noninformative Priors

Let  $X_1 = (X_{11}, \ldots, X_{1n_1})$  be a random sample of size  $n_1$  from a lognormal population with parameters  $\mu_1$  and  $\sigma^2$  and  $X_2 = (X_{21}, \ldots, X_{2n_2})$  be a random sample of size  $n_2$  from a lognormal population with parameters  $\mu_2$  and  $\sigma^2$ . That is,  $\log X_{1i}$  and  $\log X_{2j}$  are independently and normally distributed with means  $\mu_1$  and  $\mu_2$  respectively and common

variance  $\sigma^2$ . The parameter  $\theta_1 = \mu_1 - \mu_2$  is of interest. Note that

$$\frac{\mathrm{E}(X_{1i})}{\mathrm{E}(X_{2j})} = \frac{\exp(\mu_1 + \frac{1}{2}\sigma^2)}{\exp(\mu_2 + \frac{1}{2}\sigma^2)} = \exp(\theta_1). \tag{2.1}$$

Thus the problem for the ratio of two means is equivalent to  $\theta_1$ .

The pdf of  $x_1$ ,  $x_2$  is given by

$$f(\boldsymbol{x}_1, \boldsymbol{x}_2 : \mu_1, \mu_2, \sigma^2) \propto (\sigma^2)^{-\frac{n_1 + n_2}{2}} \exp \left\{ -\frac{1}{2\sigma^2} \sum_{i=1}^2 \sum_{j=1}^{n_i} (\log x_{ij} - \mu_i)^2 \right\}.$$
 (2.2)

In order to find priors, it is convenient to introduce orthogonal parametrization (Cox and Reid, 1987). To this end, let

$$\theta_1 = \mu_1 - \mu_2, \ \theta_2 = \frac{n_1}{n_1 + n_2} \mu_1 + \frac{n_2}{n_1 + n_2} \mu_2 \text{ and } \theta_3 = \sigma^2.$$
 (2.3)

With this parametrization, the likelihood has the alternate representation.

$$L(\theta_1, \theta_2, \theta_3) \propto \theta_3^{-\frac{n_1 + n_2}{2}} \exp \left[ -\frac{1}{2\theta_3} \left\{ \sum_{j=1}^{n_1} (\log x_{1j} - \frac{n_2}{n_1 + n_2} \theta_1 - \theta_2)^2 + \sum_{j=1}^{n_2} (\log x_{2j} + \frac{n_1}{n_1 + n_2} \theta_1 - \theta_2)^2 \right\} \right].$$

$$(2.4)$$

Based on (2.3), the Fisher information matrix is given by

$$I(\theta) = \text{Diag}\left(\frac{n_1 n_2}{(n_1 + n_2)\theta_3}, \frac{n_1 + n_2}{\theta_3}, \frac{n_1 + n_2}{2\theta_3^2}\right).$$
 (2.5)

Thus  $\theta_1$  is orthogonal to  $(\theta_2, \theta_3)$  in the sense of Cox and Reid (1987).

We begin with Jeffreys' prior given by

$$\pi_J(\boldsymbol{\theta}) \propto |\boldsymbol{I}(\boldsymbol{\theta})|^{\frac{1}{2}}$$

$$\propto \theta_3^{-2}.$$
(2.6)

This is a reference prior when all parameters are treated as equally important.

We consider now other reference priors. Let  $\boldsymbol{\theta}_{(1)} = \{\theta_1\}$  and  $\boldsymbol{\theta}_{(2)} = \{\theta_2, \theta_3\}$ . Then the Fisher information matrix can be expressed as

$$I(\boldsymbol{\theta}) = \mathrm{Diag}(h_1(\boldsymbol{\theta}), h_2(\boldsymbol{\theta})),$$

where  $h_1(\boldsymbol{\theta}) = n_1 n_2 / ((n_1 + n_2)\theta_3)$ ,  $h_2(\boldsymbol{\theta}) = \text{Diag}((n_1 + n_2)/\theta_3, (n_1 + n_2)/2\theta_3^2)$ . Hence, from Theorem 1 of Datta and Ghosh (1995c) the two-group reference prior is given by

$$\pi_R(\boldsymbol{\theta}) \propto \theta_3^{-\frac{3}{2}}.\tag{2.7}$$

The one-at-a-time reference prior with the partition  $\{\theta_1\}$ ,  $\{\theta_2\}$ ,  $\{\theta_3\}$  where  $\theta_1$  is the parameter of interest, while the remaining parameters are considered in any arbitrary order of importance. Writing the Fisher information matrix as

$$I(\boldsymbol{\theta}) = \operatorname{Diag}(h_1(\boldsymbol{\theta}), h_2(\boldsymbol{\theta}), h_3(\boldsymbol{\theta})),$$

where  $h_1(\boldsymbol{\theta})$  is same as before, but  $h_2(\boldsymbol{\theta}) = (n_1 + n_2)/\theta_3$  and  $h_3(\boldsymbol{\theta}) = (n_1 + n_2)/(2\theta_3^2)$ . Once again, an appeal to Theorem 1 of Datta and Ghosh (1995c) leads to the one-at-a-time reference prior as

$$\pi_R^*(\boldsymbol{\theta}) \propto \theta_3^{-1}. \tag{2.8}$$

Let  $X = (X_1, X_2)$ . For a prior  $\pi$ , let  $\theta_1^{1-\alpha}(\pi; X)$  denote the  $(1-\alpha)^{th}$  percentile of the posterior distribution of  $\theta_1$ , that is,

$$P^{\pi}\left[\theta_{1} \leq \theta_{1}^{1-\alpha}(\pi; \boldsymbol{X}) | \boldsymbol{X}\right] = 1 - \alpha, \tag{2.9}$$

where  $\boldsymbol{\theta} = (\theta_1, \dots, \theta_t)^T$  and  $\theta_1$  is the parameter of interest. We want to find priors  $\pi$  for which

$$P[\theta_1 \le \theta_1^{1-\alpha}(\pi; \boldsymbol{X}) | \boldsymbol{\theta}] = 1 - \alpha + o(n^{-u})$$
(2.10)

for some u > 0, as n goes to infinity. Priors  $\pi$  satisfying (2.10) are called matching priors. If u = 1/2, then  $\pi$  is referred to as a first order matching prior, while if u = 1,  $\pi$  is referred to as a second order matching prior.

From Tibshirani (1989), the class of first order probability matching priors is characterized by

$$\pi_F(\boldsymbol{\theta}) \propto \boldsymbol{I}_{\theta_1 \theta_1}(\boldsymbol{\theta})^{\frac{1}{2}} q(\theta_2, \theta_3)$$
$$\propto \theta_3^{-\frac{1}{2}} q(\theta_2, \theta_3), \tag{2.11}$$

where q is any arbitrary positive-valued function differentiable in its arguments. Jeffreys' prior as well as the other reference priors are all first order probability matching priors.

Clearly the class of prior given in (2.11) is quite large and it is important to narrow down this class of priors. To this end, we consider the class of second order probability matching priors as given Mukerjee and Ghosh (1997). Due to the orthogonality of  $\theta_1$  with  $(\theta_2, \theta_3)$ , the class of second order probability matching priors is characterized by solving a partial differential equation

$$\frac{1}{6}q(\theta_2, \theta_3)\frac{\partial}{\partial \theta_1} \left(I_{11}^{-\frac{3}{2}}L_{1,1,1}\right) + \sum_{\nu=2}^{3} \sum_{s=2}^{3} \frac{\partial}{\partial \theta_{\nu}} \left\{I_{11}^{-\frac{1}{2}}L_{11s}I^{s\nu}q(\theta_2, \theta_3)\right\} = 0, \tag{2.12}$$

where  $L_{1,1,1} = \mathbb{E}\Big[\big(\partial \log L/\partial \theta_1\big)^3\Big]$ ,  $L_{11s} = \mathbb{E}\Big[\partial^3 \log L/\partial \theta_1^2 \partial \theta_s\Big]$ , s = 2, 3 and  $I^{s\nu}$  is the  $(s,\nu)^{th}$  element of  $I^{-1}(\theta)$ , the inverse of the Fisher information matrix. Then (2.12) simplifies to

$$\frac{\partial}{\partial \theta_3} \left\{ \theta_3^{\frac{1}{2}} q(\theta_2, \theta_3) \right\} = 0. \tag{2.13}$$

A general class of solutions to the above equation is given by

$$q(\theta_2, \theta_3) \propto \theta_3^{-\frac{1}{2}} g(\theta_2), \tag{2.14}$$

where g is a positive function, differential in its argument, but is otherwise arbitrary. Thus the resulting second order probability matching prior is

$$\pi_S(\boldsymbol{\theta}) \propto \theta_3^{-1} g(\theta_2).$$
 (2.15)

We consider a particular second order matching prior when g is a constant in the above matching prior. This prior is given by  $\theta_3^{-1}$  which is the same as  $\pi_R^*(\theta)$ .

Based on the above calculations, it follows that  $\pi_J(\boldsymbol{\theta})$ ,  $\pi_R(\boldsymbol{\theta})$  and  $\pi_S(\boldsymbol{\theta})$  belong to a general class of priors of the form

$$\pi_a(\boldsymbol{\theta}) \propto \theta_3^{-a}. \tag{2.16}$$

In particular, the choices a=2, 3/2 and 1 lead respectively to  $\pi_J$ ,  $\pi_R$  and  $\pi_S$ . In the original parameterization, the prior  $\pi_a(\theta)$  reduces to

$$\pi_a(\mu_1, \mu_2, \sigma^2) \propto (\sigma^2)^{-a}$$
. (2.17)

We investigate the propriety of posteriors for a general class of priors which include the Jeffreys' prior, the reference prior and the second order matching prior. We consider the class of priors

$$\pi(\boldsymbol{\theta}) \propto \theta_3^{-a},\tag{2.18}$$

where a > 0. The following general theorem provides the propriety of posteriors.

**Theorem 2.1** The posterior distribution of  $\theta$  under the prior  $\pi$ , (2.18), is proper if  $n_1 + n_2 + 2a - 3 > 0$ .

The proof of this theorem is straightforward and is omitted. The marginal posterior of  $\theta_1$  under the prior (2.18) is given by

$$\pi(\theta_1|\boldsymbol{x}_1,\boldsymbol{x}_2) \propto \left\{ S^2 + \frac{n_1 n_2}{n_1 + n_2} (\theta_1 - \bar{y}_1 + \bar{y}_2)^2 \right\}^{-\frac{n_1 + n_2 + 2\alpha - 3}{2}},$$
 (2.19)

where  $y_{ij} = \log x_{ij}$ ,  $\bar{y}_i = 1/n_i \sum_{j=1}^{n_i} y_{ij}$  and  $S^2 = \sum_{i=1}^2 \sum_{j=1}^{n_i} (y_{ij} - \bar{y}_i)^2$ ,  $j = 1, \ldots, n_i$ , i = 1, 2.

### 3. Simulation Study

We evaluate the frequentist coverage probability by investigating the credible interval of the marginal posteriors density of  $\theta_1$  under the noninformative prior  $\pi$  given in (2.18) for several configurations  $(\mu_1, \mu_2, \sigma^2)$  and  $(n_1, n_2)$ . That is to say, the frequentist coverage of a  $(1-\alpha)^{th}$  posterior quantile should be close to  $1-\alpha$ .

			0.05			0.95		
$n_1$	$n_2$	$\sigma^2$	$\pi_J$	$\pi_R$	$\pi_S$	$\pi_J$	$\pi_R$	$\pi_S$
5	5	0.5	0.073	0.062	0.050	0.929	0.940	0.950
		1	0.072	0.061	0.051	0.929	0.940	0.950
		10	0.073	0.063	0.051	0.928	0.940	0.951
5	10	0.5	0.061	0.054	0.048	0.940	0.945	0.951
		1	0.064	0.058	0.051	0.936	0.942	0.951
		10	0.061	0.055	0.049	0.935	0.943	0.949
10	10	0.5	0.063	0.057	0.053	0.941	0.945	0.952
		1	0.060	0.056	0.049	0.939	0.944	0.949
		10	0.059	0.055	0.050	0.936	0.941	0.946
10	20	0.5	0.056	0.052	0.050	0.945	0.948	0.951
		1	0.056	0.053	0.050	0.943	0.945	0.949
		10	0.056	0.052	0.048	0.946	0.948	0.951

Table 3.1: Comparisons of Coverage Probability for Posterior Quantiles of  $\theta_1$  with Equal Variances

For fixed  $(\mu_1, \mu_2, \sigma^2)$  and  $(n_1, n_2)$ , we take 10,000 independent random samples of  $(\mathbf{x}_1, \mathbf{x}_2)$  from the model (2.2).

The Table 3.1 gives numerical values of the frequentist coverage probabilities of 0 to 0.05 and 0 to 0.95 posterior quantiles under the Jeffreys', reference prior and second order matching prior.

In Table 3.1, we can observe that the second order matching prior  $\pi_S$  meets very well the target coverage probabilities. This is intuitively clear since  $\pi_S$  is a second order matching prior, but  $\pi_J$ ,  $\pi_R$  are not. Also note that the results of table are not much sensitive to the change of the values of  $\sigma^2$ . Thus we recommend to use the second order matching prior for the Bayesian inference for the ratio of the lognormal means with equal variances.

### 4. Numerical Example in Bioequivalence Problems

In this section, we will illustrate our Bayesian method using a real-life example. AUC data from an experiment with simultaneous administration of a test formulation  $(x_1)$  and a reference formulation  $(x_2)$  in six male subjects (Alkalay et al., 1980). The AUC data from this study are presented in Table 4.1  $(n_1 = n_2 = 6)$ . The sample means are 38.618 and 39.440, and the sample standard deviations are 25.112 and 26.394. After the log-transformation, the sample means are 3.499 and 3.508 and the sample standard deviations are 0.590 and 0.614.

The Shapiro-Wilk tests for the normality on the log-transformed data give a p-value of 0.191 for the test formulation group and a p-value of 0.185 for the reference formulation group, while the same tests on the original data give a p-value of 0.079 for the test

Table 4.1: AUC Data from an Experiment

$x_1$	19.96	20.37	52.42	21.37	82.98	34.61
$x_2$	19.91			19.73	85.33	34.88

formulation group and a p-value of 0.084 for the reference formulation group. Therefore, the log-transformation normalizes the data. The F-test for equal variances of the log-transformed data between the two groups gives a p-value of 0.932, and therefore the log-transformation stabilizes the variances.

We have two theories. Theory one says that two formulations are not bioequivalence, i.e.,  $H_0: \theta_1 \leq \kappa_L$  or  $\theta_1 \geq \kappa_U$ . Theory two says that two formulations are bioequivalence, i.e.,  $H_a: \kappa_L < \theta_1 < \kappa_U$  (Berger and Hsu, 1996). Here  $\theta_1$  is that logarithm transformation be taken the ratio of the lognormal means.

The posterior probability of the null hypothesis is

$$P(H_0|\boldsymbol{x}_1, \boldsymbol{x}_2) = \frac{\int_{-\infty}^{\kappa_L} \pi(\theta_1|\boldsymbol{x}_1, \boldsymbol{x}_2) d\theta_1 + \int_{\kappa_U}^{\infty} \pi(\theta_1|\boldsymbol{x}_1, \boldsymbol{x}_2) d\theta_1}{\int_{-\infty}^{\infty} \pi(\theta_1|\boldsymbol{x}_1, \boldsymbol{x}_2) d\theta_1},$$
(4.1)

and the posterior probability of the alternative hypothesis is

$$P(H_a|\mathbf{x}_1, \mathbf{x}_2) = \frac{\int_{\kappa_L}^{\kappa_U} \pi(\theta_1|\mathbf{x}_1, \mathbf{x}_2) d\theta_1}{\int_{-\infty}^{\infty} \pi(\theta_1|\mathbf{x}_1, \mathbf{x}_2) d\theta_1},$$
(4.2)

where  $\pi(\theta_1|\boldsymbol{x}_1,\boldsymbol{x}_2)$  is the marginal posterior of  $\theta_1$  is given in (2.19).

In this case,  $n_1 = 6$ ,  $n_2 = 6$ ,  $\bar{y}_1 = 3.499$ ,  $\bar{y}_2 = 3.508$  and  $S^2 = 3.627$ .

Table 4.2 provides the posterior probabilities of the null hypothesis  $H_0$  and alternative hypothesis  $H_a$  corresponding to Jeffreys' prior  $(\pi_J)$ , two group reference prior  $(\pi_R)$  and second order probability matching prior  $(\pi_S)$ . Also Table 4.2 provides the Bayes factors for  $H_0$  versus  $H_a$ .

If we use  $\kappa_U = -\kappa_L = \log 1.25$ , the Bayes factor gives the weak evidence for  $H_a$  for  $\pi_J$ . Hence two formulations are bioequivalence. But the Bayes factor gives the weak evidence for  $H_0$  for  $\pi_R$  and  $\pi_S$ . Hence two formulations are bioinequivalence. This discrepancy in the conclusion may be due to the poor performance of  $\pi_J$  as we already noticed in Section 3.

Table 4.2: Posterior Probabilities and Bayes Factors with Equal Variances

	$\kappa_U = -\kappa_L = \log 1.25$				
	$\pi_J$	$\pi_R$	$\pi_S$		
$P(H_0 oldsymbol{x}_1,oldsymbol{x}_2)$	0.496	0.515	0.536		
$P(H_0 m{x}_1,m{x}_2) \ P(H_a m{x}_1,m{x}_2) \ B_{12}$	0.504	0.485	0.464		
$B_{12}$	0.984	1.062	1.155		

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