Notes on Bayesian Prevalence

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Introduction

We consider a population of units (participants or spike-sorted single neuron spike trains) which are of two types. Within the population, a proportion γ possess some definable effect, while the proportion of units in the population who do not possess this effect is $1-\gamma$. The *prevalence* of the defined effect within the population is γ , ($0<\gamma<1$). A random sample of n units is selected from the population and each unit undergoes a test procedure, in which the presence of the defined effect is investigated using a significance test. It is assumed that for each unit the significance level of the test is a (1 - specificity) and the power of the test (sensitivity) is b. Thus, the probability that a randomly selected unit from the population who does not possess the defined effect will produce a significant result is a, while the probability that a randomly selected unit from the population who does possess the defined effect will produce a significant result is b.

A binary variable – shows a significant effect or does not show a significant effect is recorded for each unit in the sample, and we suppose that the total number of units who show a significant effect, out of the n tested, is k. Let θ be the probability that a randomly selected unit from the population will show a significant effect. Then

$$\theta = (1 - \gamma)a + \gamma b = a + (b - a)\gamma. \tag{1}$$

We will develop the modelling in terms of the parameter θ , and later use (1) to find appropriate results in terms of the prevalence, γ .

Modelling

Assuming that the test results on the performance of the units are independent and that the parameter θ is the same for all units in the population. Let the random variable X denote the number of units out of the n tested which show a significant effect at significance level a. Then X follows a binomial distribution and

$$\Pr(X = k | \theta) = \binom{n}{k} \theta^k (1 - \theta)^{n-k}, \quad k = 0, 1, \dots, n, \quad (0 < \theta < 1).$$
 (2)

We now define a prior distribution to characterise the prior uncertainty about θ . First, we note that under the uncontroversial assumption that b > a, we find from (1) that $\theta > a$. Also, since

 γ < 1, we find that θ < b. The claim regarding the assumption that b > a is perfectly reasonable since it would make no sense to employ a test procedure for which the power is less than the significance level. It follows that a < θ < b.

The conjugate prior for θ is the beta distribution so, bearing in mind the constraint on θ , we assume that the prior distribution for θ is the following truncated beta distribution with probability density function

$$p(\theta|a,b,r,s) = \frac{1}{B(r,s)} \frac{\theta^{r-1}(1-\theta)^{s-1}}{[F(b;r,s) - F(a;r,s)]}, \quad a < \theta < b, \quad (r > 0, s > 0),$$

$$\equiv \frac{\text{Beta}(r,s)}{[F(b;r,s) - F(a;r,s)]}$$
(3)

where F(x;r,s) is the cumulative distribution function (cdf) of θ given by the following beta cdf,

$$F(x;r,s) = \frac{1}{B(r,s)} \int_0^x \theta^{r-1} (1-\theta)^{s-1} dt$$
 (4)

Beta(r, s) is the pdf of the beta distribution and B(r,s) is the beta function, both having parameters r, s. The selection of values for the parameters r, s depends on prior information about θ . In the absence of any prior information about θ we will use the choice r=1, s=1 in practical applications, while keeping the notation general in the formulation. This corresponds to the a priori assumption that the prior uncertainty regarding θ can be represented by a uniform distribution on the interval (a,b).

We define $m_1 \equiv k + r, m_2 \equiv n - k + s$. Combination of the likelihood in (2) with the prior in (3) by means of Bayes' theorem gives the posterior probability density function for θ as

$$p(\theta|k,a,b,r,s) \propto \theta^{m_1-1}(1-\theta)^{m_2-1}, \quad a < \theta < b,$$

and so the posterior p.d.f. is the truncated beta distribution

$$p(\theta|k,a,b,r,s) = \frac{\text{Beta}(m_1, m_2)}{[F(b; m_1, m_2) - F(a; m_1, m_2)]}, \quad a < \theta < b.$$
 (5)

In the sequel, the cdf and its inverse - the quantile function - for the truncated beta distribution will be required so we now provide expression for these functions: C(x) for the cdf and Q(p) for the quantile function.

$$C(x) = \Pr(\theta < x) = \int_{a}^{x} p(\theta | k, a, b, r, s) d\theta = \frac{F(x; m_1, m_2) - F(a; m_1, m_2)}{F(b; m_1 m_2) - F(a; m_1, m_2)}.$$
 (6)

Suppose that we wish to find the pth quantile, x, of the truncated beta distribution in (5). That is: we wish to solve the equation

$$C(x) \equiv \int_{a}^{x} p(\theta|k, a, b, r, s) d\theta = p.$$
 (7)

Then using (6), we may write this equation as

$$F(x; m_1, m_2) = (1 - p)F(a; m_1, m_2) + pF(b; m_1, m_2)$$
(8)

and so

$$x = F^{-1}[(1-p)F(a; m_1, m_2) + pF(b; m_1, m_2)].$$

Thus, the quantile function for the pth quantile of the truncated beta distribution is

$$Q(p) = F^{-1} \left[(1 - p)F(a; m_1, m_2) + pF(b; m_1 m_2) \right], \tag{9}$$

where as before F is the cdf of the beta distribution given in (4).

Applications

We now derive some applications of the truncated beta distribution from (5) in relation to the prevalence, γ .

Posterior distribution of γ

Using the standard result for transforming random variables, we find that the posterior p.d.f. for γ is

$$p(\gamma|k,a,b,r,s) = c[a + (b-a)\gamma]^{m_1-1}[1 - a - (b-a)\gamma]^{m_2-1}, \quad (0 < \gamma < 1), \tag{10}$$

where the constant c has the form

$$c = \frac{b - a}{B(m_1, m_2)[F(b; m_1, m_2) - F(a; m_1, m_2)]}.$$
 (11)

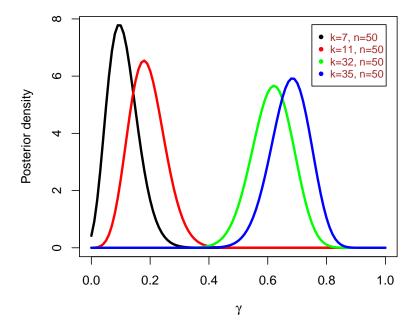


Figure 1: Posterior pdfs of population prevalence for four different choices of the values of (k, n), where k is the number of units, out of n tested, which show a significant result.

Figure 1 shows some posterior pdfs for γ .

Lower bound for γ

We can determine a lower bound , γ_c , for the prevalence by exploiting the relationship between θ and γ from (1) in the form

$$\gamma = \frac{\theta - a}{b - a} \tag{12}$$

and we note that

$$\gamma \ge \gamma_c \iff \theta \ge \theta_c \equiv a + (b - a)\gamma_c.$$
 (13)

Then from (1) and (7),

$$\Pr(\gamma \ge \gamma_c) = \Pr(\theta \ge \theta_c) = 1 - \Pr(\theta < \theta_c) \equiv 1 - C(\theta_c).$$

Using (5) we first find a posterior interval for θ of the form $(\theta_c, 1)$ which has posterior probability p by solving

$$\int_{\theta_c}^b p(\theta|k,a,b,r,s) d\theta = p,$$

which can be written as

$$C(\theta_c) = 1 - p,$$

so that from (9)

$$\theta_c = Q(1-p)$$
.

Then from (13) we find the corresponding lower bound for γ as

$$\gamma_c = \frac{\theta_c - a}{b - a} \tag{14}$$

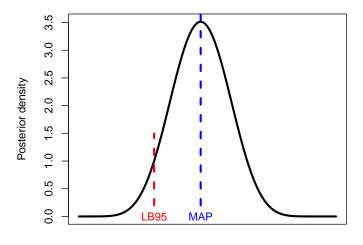


Figure 2: An illustration of the 0.95 lower bound for γ , denoted by LB95, as well as the MAP estimate of γ , when k = 10 units, out of a total of n units, show a significant result.

γ

MAP estimate of γ

The MAP estimate is the posterior mode for γ . It is given by

$$\begin{cases} 0 & \hat{\theta} \le a \\ \frac{\hat{\theta} - a}{b - a} & a < \hat{\theta} < b, \text{ where } \hat{\theta} = \frac{m_1 - 1}{m_1 + m_2 - 2} \\ 1 & \hat{\theta} \ge b \end{cases}$$

When r = 1, s = 1, $\hat{\theta} = k/n$.

An illustration of a 0.95 lower bound as well as a MAP estimate are shown in Figure 2.

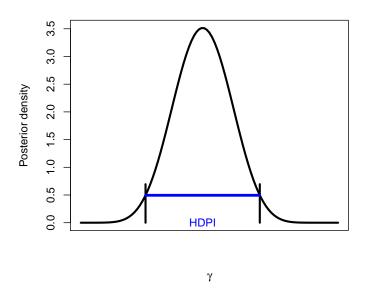


Figure 3: The HPDI when k = 10 and n = 20.

Highest posterior density interval for γ

Depending on the shape of the posterior pdf for γ , the HPDI can take several forms. It could be (i) a two-sided interval, (ii) a one-sided interval or (iii) a set of disjoint intervals. Case (i) happen when the posterior pdf is unimodal and the mode occurs when γ is neither 0 nor 1. Case (ii) occurs when posterior mode occurs when $\gamma = 0$ or when $\gamma = 1$. Case (iii) is not relevant here but it occurs when the posterior pdf is multimodal. We focus on Case (i). Then the HPDI with posterior probability p is the shortest interval of values of γ for which the posterior probability that γ lies between the endpoints of this interval is equal to p. We assume that r = 1, s = 1.

We first find the HPDI for θ which has posterior probability p, and then use relation (12) to derive the corresponding interval for γ . Mathematically, it is required to find endpoints e_1 , e_2 for θ such that

$$C(e_2) - C(e_1) = p,$$

 $p(e_2) - p(e_1) = 0,$

where C is the cdf of the truncated beta distribution defined in (7) and p(e) is the posterior pdf for

 θ in (5) evaluated at $\theta = e$, with r = 1, s = 1.

The HPDI for γ is then computed using (12). One-sided intervals occur when k=0 or k=n or if the HPDI for θ has a left-hand endpoint less than or equal to a or a right-hand endpoint that is greater than or equal to b. An illustration is shown in Figure 3.

Sampling distribution

For a given unknown population prevalence γ , there will be variation in the number k of significant results obtained from repeated sets of tests in which n units are tested. Various statistics, such as (i) the length of the HDPI for γ (ii) the MAP estimate of γ and (iii) a lower bound for γ , are all subject to this sampling variation. It is useful then to consider the sampling distribution of each of these statistics and then compute its mean and standard deviation.

For a given number n of units, the value of k can be anything from 0 to n, with probability distribution given in (2). Let S be a statistic of interest which has value s_k when k out of n tests are significant (k = 0, 1, ..., n). Then the mean value of S is

$$\mu_n = \sum_{k=0}^n \Pr(X = k | \theta) s_k \tag{15}$$

and the standard deviation of *S* is

$$\sigma_n = \sqrt{\sum_{k=0}^n \Pr(X = k | \theta) (s_k - \mu_n)^2}.$$
(16)

Formulae (15), (16) are then applied by taking S in turn to be (i) the length of the HPDI for γ , (ii) the MAP estimate of γ and (iii) a lower bound for γ , or any other relevant statistic.