# 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 $\gamma$ 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 $\gamma,(0<\gamma<1)$. 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 $\theta$ be the probability that a randomly selected unit from the population will show a significant effect. Then

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

We will develop the modelling in terms of the parameter $\theta$, and later use (1) to find appropriate results in terms of the prevalence, $\gamma$.

## Modelling

Assuming that the test results on the performance of the units are independent and that the parameter $\theta$ 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

$$
\begin{equation*}
\operatorname{Pr}(X=k \mid \theta)=\binom{n}{k} \theta^{k}(1-\theta)^{n-k}, \quad k=0,1, \ldots, n, \quad(0<\theta<1) . \tag{2}
\end{equation*}
$$

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

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

$$
\begin{align*}
p(\theta \mid a, b, r, s) & =\frac{1}{B(r, s)} \frac{\theta^{r-1}(1-\theta)^{s-1}}{[F(b ; r, s)-F(a ; r, s)]}, \quad a<\theta<b, \quad(r>0, s>0)  \tag{3}\\
& \equiv \frac{\operatorname{Beta}(r, s)}{[F(b ; r, s)-F(a ; r, s)]}
\end{align*}
$$

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

$$
\begin{equation*}
F(x ; r, s)=\frac{1}{B(r, s)} \int_{0}^{x} \theta^{r-1}(1-\theta)^{s-1} d t \tag{4}
\end{equation*}
$$

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 $\theta$. In the absence of any prior information about $\theta$ 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 $\theta$ 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 $\theta$ as

$$
p(\theta \mid k, a, b, r, s) \propto \theta^{m_{1}-1}(1-\theta)^{m_{2}-1}, \quad a<\theta<b
$$

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

$$
\begin{equation*}
p(\theta \mid k, a, b, r, s)=\frac{\operatorname{Beta}\left(m_{1}, m_{2}\right)}{\left[F\left(b ; m_{1}, m_{2}\right)-F\left(a ; m_{1}, m_{2}\right)\right]}, \quad a<\theta<b . \tag{5}
\end{equation*}
$$

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 $c d f$ and $Q(p)$ for the quantile function.

$$
\begin{equation*}
C(x)=\operatorname{Pr}(\theta<x)=\int_{a}^{x} p(\theta \mid k, a, b, r, s) d \theta=\frac{F\left(x ; m_{1}, m_{2}\right)-F\left(a ; m_{1}, m_{2}\right)}{F\left(b ; m_{1} m_{2}\right)-F\left(a ; m_{1}, m_{2}\right)} . \tag{6}
\end{equation*}
$$

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

$$
\begin{equation*}
C(x) \equiv \int_{a}^{x} p(\theta \mid k, a, b, r, s) d \theta=p . \tag{7}
\end{equation*}
$$

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

$$
\begin{equation*}
F\left(x ; m_{1}, m_{2}\right)=(1-p) F\left(a ; m_{1}, m_{2}\right)+p F\left(b ; m_{1}, m_{2}\right) \tag{8}
\end{equation*}
$$

and so

$$
x=F^{-1}\left[(1-p) F\left(a ; m_{1}, m_{2}\right)+p F\left(b ; m_{1}, m_{2}\right)\right] .
$$

Thus, the quantile function for the $p$ th quantile of the truncated beta distribution is

$$
\begin{equation*}
Q(p)=F^{-1}\left[(1-p) F\left(a ; m_{1}, m_{2}\right)+p F\left(b ; m_{1} m_{2}\right)\right], \tag{9}
\end{equation*}
$$

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, $\gamma$.

## Posterior distribution of $\gamma$

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

$$
\begin{equation*}
p(\gamma \mid k, a, b, r, s)=c[a+(b-a) \gamma]^{m_{1}-1}[1-a-(b-a) \gamma]^{m_{2}-1}, \quad(0<\gamma<1), \tag{10}
\end{equation*}
$$

where the constant $c$ has the form

$$
\begin{equation*}
c=\frac{b-a}{B\left(m_{1}, m_{2}\right)\left[F\left(b ; m_{1}, m_{2}\right)-F\left(a ; m_{1}, m_{2}\right)\right]} . \tag{11}
\end{equation*}
$$



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 $\gamma$.

## Lower bound for $\gamma$

We can determine a lower bound, $\gamma_{c}$, for the prevalence by exploiting the relationship between $\theta$ and $\gamma$ from (1) in the form

$$
\begin{equation*}
\gamma=\frac{\theta-a}{b-a} \tag{12}
\end{equation*}
$$

and we note that

$$
\begin{equation*}
\gamma \geq \gamma_{c} \Longleftrightarrow \theta \geq \theta_{c} \equiv a+(b-a) \gamma_{c} . \tag{13}
\end{equation*}
$$

Then from (1) and (7),

$$
\operatorname{Pr}\left(\gamma \geq \gamma_{c}\right)=\operatorname{Pr}\left(\theta \geq \theta_{c}\right)=1-\operatorname{Pr}\left(\theta<\theta_{c}\right) \equiv 1-C\left(\theta_{c}\right) .
$$

Using (5) we first find a posterior interval for $\theta$ of the form $\left(\theta_{c}, 1\right)$ which has posterior probability $p$ by solving

$$
\int_{\theta_{c}}^{b} p(\theta \mid k, a, b, r, s) d \theta=p,
$$

which can be written as

$$
C\left(\theta_{c}\right)=1-p,
$$

so that from (9)

$$
\theta_{c}=Q(1-p) .
$$

Then from (13) we find the corresponding lower bound for $\gamma$ as

$$
\begin{equation*}
\gamma_{c}=\frac{\theta_{c}-a}{b-a} \tag{14}
\end{equation*}
$$


$\gamma$

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

## MAP estimate of $\gamma$

The MAP estimate is the posterior mode for $\gamma$. It is given by

$$
\begin{cases}0 & \hat{\theta} \leq a \\ \hat{\theta}-a \\ b-a & a<\hat{\theta}<b, \quad \text { where } \quad \hat{\theta}=\frac{m_{1}-1}{m_{1}+m_{2}-2} \\ 1 & \hat{\theta} \geq b\end{cases}
$$

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.

$\gamma$

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

## Highest posterior density interval for $\gamma$

Depending on the shape of the posterior pdf for $\gamma$, 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 $\gamma$ 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 $\gamma$ for which the posterior probability that $\gamma$ lies between the endpoints of this interval is equal to $p$. We assume that $r=1, s=1$.

We first find the HPDI for $\theta$ which has posterior probability $p$, and then use relation (12) to derive the corresponding interval for $\gamma$. Mathematically, it is required to find endpoints $e_{1}, e_{2}$ for $\theta$ such that

$$
\begin{aligned}
C\left(e_{2}\right)-C\left(e_{1}\right) & =p, \\
p\left(e_{2}\right)-p\left(e_{1}\right) & =0,
\end{aligned}
$$

where $C$ is the $c d f$ of the truncated beta distribution defined in (7) and $p(e)$ is the posterior pdf for
$\theta$ in (5) evaluated at $\theta=e$, with $r=1, s=1$.
The HPDI for $\gamma$ is then computed using (12). One-sided intervals occur when $k=0$ or $k=n$ or if the HPDI for $\theta$ 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 $\gamma$, 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 $\gamma$ (ii) the MAP estimate of $\gamma$ and (iii) a lower bound for $\gamma$, 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 $(k=0,1, \ldots, n)$. Then the mean value of $S$ is

$$
\begin{equation*}
\mu_{n}=\sum_{k=0}^{n} \operatorname{Pr}(X=k \mid \theta) s_{k} \tag{15}
\end{equation*}
$$

and the standard deviation of $S$ is

$$
\begin{equation*}
\sigma_{n}=\sqrt{\sum_{k=0}^{n} \operatorname{Pr}(X=k \mid \theta)\left(s_{k}-\mu_{n}\right)^{2}} \tag{16}
\end{equation*}
$$

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

