



Binomial distribution and estimation of the true prevalence of infected animals when pooled samples are used

Dear Sir,

A binomial experiment is one that possesses the following properties:

1. The experiment consists of several identical trials.
2. Each trial results in 1 of 2 outcomes; one, a success (negative for contamination), the other, a failure (positive for contamination).
3. The probability of success on a single trial is equal to p and remains the same from trial to trial. The probability of a failure is equal to $q = 1 - p$.
4. The trials are independent.

The binomial theorem can be most easily seen by observing the pattern obtained in calculating $(p + q)^n$ for $n = 1, 2, 3, \dots$. In a general terms, the following formula is called the binomial theorem:

$$(p + q)^n = p^n + n p^{n-1} q + [n(n-1)]/n! p^{n-2} q^2 + \dots + q^n$$

Example: The published data of Sorensen et al (1). Feces were cultured for the presence of *Mycobacterium paratuberculosis*. Culture was not carried out on individual fecal samples. Approximately 2 g from each of 3 individual 60-mL fecal samples (a total of approximately 6 g) were pooled and mixed. Samples were pooled in the order of sampling. Five hundred fecal pools were tested for *M. paratuberculosis*. Thirty (6.0%) of the 500 fecal pools became overgrown with fungus. No determination could be made with these cultures and they were removed from the subsequent analysis. Sixteen of the 470 readable cultures (3.4%) showed growth of *M. paratuberculosis* (positive) and 454 cultures did not show growth of *M. paratuberculosis* (negative). If we assume that the prevalence of positive and negative fecal pools was q and p , respectively, by using binomial distribution, it could be concluded that:

- p^3 is equal to frequency of fecal pools that 3 animals were not infected with *M. paratuberculosis* (negative).
- $3p^2q$ is equal to frequency of fecal pools that 2 animals were negative and 1 was positive.

- $3pq^2$ is equal to frequency of fecal pools that 1 animal was negative and 2 were positive.
- q^3 is equal to frequency of fecal pools that 3 animals were positive.

As mentioned above, the frequencies of 1, 2, and all 3 individual fecal samples in the positive fecal pool are not equal to each other.

By using the frequency of negative fecal pools (p^3), we can estimate the p , and then the prevalence of infected cows ($q = 1 - p$).

$$p = (p^3)^{1/3} = (\text{frequency of negative fecal pools})^{1/3} = (0.9659)^{1/3} = 0.9885 \text{ or } 98.85\%$$

$$q = \text{frequency of infected cows} = 1 - p = 0.0115 \text{ or } 1.15\%$$

The true cow-level prevalence should be estimated by using the sensitivity and specificity of the diagnostic test.

In summary, by using data from pooled samples of n animals (where $n = 2, 3, 4, \dots$), we can estimate the true prevalence of infected animals and its standard deviation (s) as follows:

$$\begin{aligned} \text{True prevalence of infected animals} &= \\ &[1 - (\text{Frequency of negative samples})^{1/n}] \\ &(\text{Specificity} - 1)/[\text{Sensitivity} + (\text{Specificity} - 1)] \\ s \text{ of true prevalence} &= [(\text{True prevalence}) \\ &(1 - \text{True prevalence})/\text{Sample size}]^{1/2} \end{aligned}$$

In conclusion, Sorensen et al over-estimated the true prevalence of infection at cow-level.

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Reference

1. Sorensen O, Rawluk S, Wu J, Manninen K, Ollis G. *Mycobacterium paratuberculosis* in dairy herds in Alberta. Can Vet J 2003;44:221-226.

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