

STA286 Lecture 28

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comparing two proportions

One variable in the dataset with 0's and 1's; another variable splitting observations into two groups.

The two populations are Bernoulli(p_1) and Bernoulli(p_2). The independent samples are X_{11}, \dots, X_{1n_1} and X_{21}, \dots, X_{2n_2}

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Formula for 95% interval:

$$(\hat{p}_1 - \hat{p}_2) \pm 1.96 \sqrt{\frac{\hat{p}_1(1 - \hat{p}_1)}{n_1} + \frac{\hat{p}_2(1 - \hat{p}_2)}{n_2}}$$

two non-robust confidence intervals

Every procedure I have explained so far is *robust* as long as the sample size is large enough (except for the prediction interval formula.)

In principle we could apply the patented procedure to estimate σ^2 with S^2 , using a χ^2 distribution.

We could also apply the patented procedure to estimate the ratio σ_1^2/σ_2^2 with S_1^2/S_2^2 using an F distribution.

But the results are well known to be non-robust, even with large sample sizes, so I cannot recommend them for use.

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But the results are well known to be non-robust, even with large sample sizes, so I cannot recommend them for use.

A modern computationally-intensive technique called *bootstrapping* is a good choice in these and other difficult situations.

chiselling your own estimators onto stone tablets

fun fact from mathematics

Suppose a twice-differentiable function $f(x)$ has a critical value at x_0 , and $g(x)$ is strictly increasing and twice-differentiable.

Then $g(f(x))$ also has a critical value at x_0 , and the sign of its second derivative at x_0 is the same as the sign of the second derivative of f at x_0 .

This can be seen by evaluating the left hand sides at x_0 :

$$\begin{aligned}(g(f(x)))' &= g'(f(x))f'(x) \\ (g(f(x)))'' &= g''(f(x))(f'(x))^2 + g'(f(x))f''(x)\end{aligned}$$

estimating a proportion, from first principles - I

Here's a simulated sequence of 0's and 1's from a Bernoulli(p) distribution. I know what (p), but you don't.

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## [1] 0 0 0 0 0 0 0 0 1 0 1
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What value of p between 0 and 1 is the *most likely* to have produce this sequence of 2 1's and 8 0's?

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The probability of getting this sample exactly is:

$$\begin{aligned} & (1 - p) \cdot (1 - p) \cdot (1 - p) \cdot (1 - p) \cdot (1 - p) \cdot (1 - p) \cdot (1 - p) \cdot p \cdot (1 - p) \cdot p \\ &= p^2(1 - p)^8 \end{aligned}$$

Let's call this function $L(p)$.

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Let's call this function $L(p)$.

We could maximize $L(p)$, but it's easier to maximize $\ell(p) = \log L(p)$

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$$0 = \frac{d}{dp} \ell(p) = \frac{d}{dp} (2 \log(p) + 8 \log(1 - p)) = \frac{2}{p} - \frac{8}{1 - p}$$

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The second derivative is negative, so this is a maximum.

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This is exactly the same as \hat{p} that was used as “obvious” from before.

“likelihood function” for Bernoulli

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$$L(p) = \prod_{i=1}^n p^{x_i} (1 - p)^{1-x_i}$$

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$$\text{Also: } \ell(p) = \log L(p) = \sum_{i=1}^n \log f(x_i; p)$$

likelihood function in general

Given a sequence of observations $\{x_1, \dots, x_n\}$ (“the data”) from a random variable X with pmf or pdf $f(x; \theta)$, a likelihood function $L(\theta) = L(x_1, \dots, x_n; \theta)$ for the parameter θ is defined as (for any positive g):

$$L(\theta) = g(\mathbf{x}) \underbrace{\prod_{i=1}^n f(x_i; \theta)}_{\text{real definition}}$$

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If X is discrete and f is a pmf, then $L(\theta)$ is literally the probability of the data given θ .

If X is continuous and f is a pdf, then $L(\theta)$ is not a probability, but it still provides a useful “index” for θ values.

likelihood as “index” in continuous case

Suppose $X \sim \text{Exp}(\lambda)$ and the data are: 1, 3, 8. A likelihood for λ is:

$$L(1, 3, 8; \lambda) = \lambda^3 e^{-\lambda(1+3+8)} = \lambda^3 e^{-12\lambda}$$

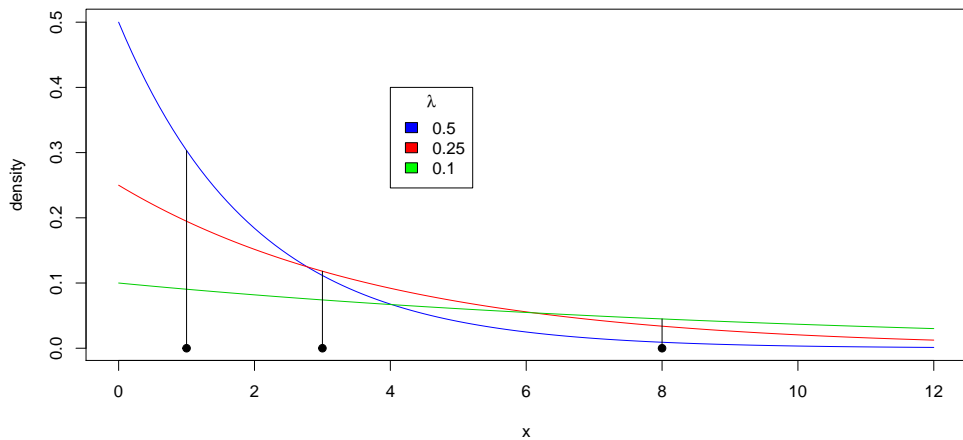
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Consider three possible candidate guesses for the true value of λ : 0.1, 0.25, and 0.5.

a possibly useless and confusing picture



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For example, suppose x_1, x_2, \dots, x_n are data observed from a $X \sim N(\mu, 1)$ population. A likelihood for μ is:

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To maximize:

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the maximum likelihood estimator

A final technicality. When you replace the data x_1, x_2, \dots, x_n with its “model”, the sample: X_1, X_2, \dots, X_n , inside the maximum likelihood estimate, you end up with the *maximum likelihood estimator*, or MLE.

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For example, the maximum likelihood estimator for μ using a sample from a $N(\mu, 1)$ population is:

$$\hat{\mu} = \bar{X}$$

Everything so far extends to vector parameters. For example (textbook example 9.21), the maximum likelihood estimates given data x_1, \dots, x_n from a $N(\mu, \sigma)$ population, the MLE for $\theta = (\mu, \sigma^2)$ are:

$$\hat{\mu} = \bar{X} \qquad \hat{\sigma}^2 = \frac{\sum_{i=1}^n (X_i - \bar{X})^2}{n}$$

properties of the maximum likelihood estimator

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3. it is “invariant”, which means $\widehat{h(\theta)} = h(\hat{\theta})$ when h is a 1-1 function.
4. it is asymptotically normal.
5. if $c\hat{\theta}$ is unbiased for some constant c , then $c\hat{\theta}$ is the unbiased estimator with the smallest variance (our “gold standard”).