Lecture 22 Confidence Interval For The Proportion

BIO210 Biostatistics

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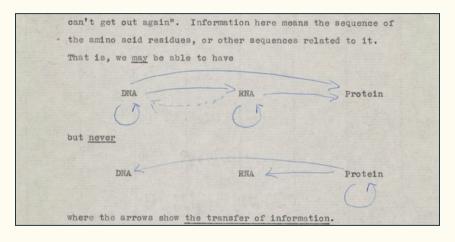
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Population Parameters We Have Learnt

Population parameters	Sample statistics
μ	\bar{x}
σ^2	s^2
σ	s
π or p	p or \hat{p}

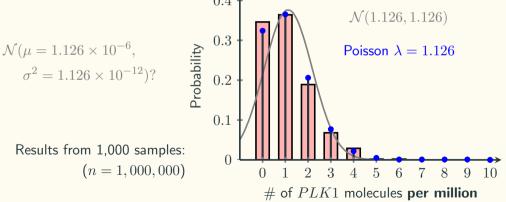
The Central Dogma



Credit: "Ideas on protein synthesis (Oct. 1956)". Wellcome Collection.

Sample Proportion Example

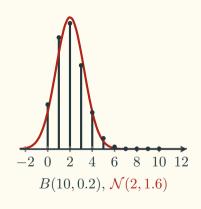
Gene expression (over-simplified RNA-seq): We know the probability of detecting PLK1 is $\pi=0.000001126088083$. If we take a random sample of n=1,000,000 mRNA molecules, what is the sampling distribution of proportion of PLK1?

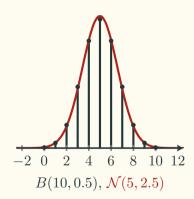


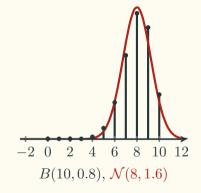
Approximation of The Binomial Distribution

$$B(n,p) \begin{cases} \dot{\sim} \ \mathcal{N}(\mu=np,\sigma^2=npq) & , \text{ when } np\geqslant 10 \text{ and } nq\geqslant 10 \\ \\ \dot{\sim} \ Pois(\lambda=np) & , \text{ when } n \text{ is large, and } p \text{ is small,} \\ \\ \text{such that } np \text{ is between } 0 \text{ and } 10. \end{cases}$$

The Limitations on np and nq







The Limitations on np and nq

- Binomial: all data are within [0, n]
- Normal: no bounds $(-\infty, +\infty)$ for data, but most are within $[\mu 3\sigma, \ \mu + 3\sigma]$
- Intuitively: when $[\mu 3\sigma, \ \mu + 3\sigma]$ is within [0, n], the approximation works well!

$$\begin{array}{lll} \mu-3\sigma>0 & \mu+3\sigma< n \\ np-3\sqrt{npq}>0 & np+3\sqrt{npq}< n \\ & np>3\sqrt{npq} & n(1-p)>3\sqrt{npq} \\ & n^2p^2>9npq & n^2q^2>9npq \\ & np>9q & nq>9p \\ & np>9(1-p)=9-9p & nq>9(1-q)=9-9q \end{array}$$

Interval Estimation For The Proportion

Goal: for a population containing an unknown proportion (π) of data of our interest, find a and b, such that $\mathbb{P}(a \leq \pi \leq b) = 0.95$.

$$\mathbb{P}\left(-1.96 \leqslant Z \leqslant 1.96\right) = 0.95$$

$$\mathbb{P}\left(-1.96 \leqslant \frac{p - \mu_P}{\sigma_P} \leqslant 1.96\right) = 0.95$$

$$\mathbb{P}\left(-1.96 \leqslant \frac{p - \pi}{\sqrt{\frac{\pi(1 - \pi)}{n}}} \leqslant 1.96\right) = 0.95$$

$$\mathbb{P}\left(p - 1.96\sqrt{\frac{\pi(1 - \pi)}{n}} \leqslant \pi \leqslant p + 1.96\sqrt{\frac{\pi(1 - \pi)}{n}}\right) = 0.95$$

Confidence Interval For The Proportion

95% CI For The Sample Proportion

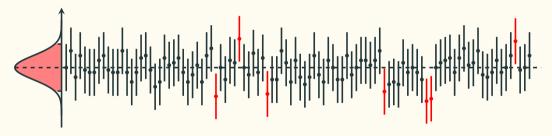
The Wald Interval:

$$\left[p - 1.96\sqrt{\frac{p(1-p)}{n}}, p + 1.96\sqrt{\frac{p(1-p)}{n}}\right]$$

• Not using t-distribution? - You don't need to! Remember $\sigma_P = \sqrt{\frac{\pi(1-\pi)}{n}}$, and when p is calculated to estimate π , then σ_P is automatically determined, unlike in the situation of the mean, where you have to do extra (independent) calculation of s to estimate σ , which causes the extra error.

Simulation of 95% CI For The Proportion

100 95% CI for the proportion, constructed using the Wald interval



An Example in Lecture 1

Probability vs. Statistics

Probability: Previous studies showed that the drug was 80% effective. Then

we can anticipate that for a study on 100 patients, on average 80 will be cured and at least 65 will be cured with 99.99%

chance.

Statistics: We observe that 78/100 patients were cured by the drug. We

will be able to conclude that we are 95% confident that for other studies the drug will be effective on between 69.88% and

86.11% of patients.

Sample Size Estimation Using Confidence Interval of The Proportion

Estimate Sample Size: We want to estimate the percentage of people cured by the drug. Suppose we could draw a truly random sample, and we want a 95% confidence interval estimation with a margin of error no more than $\pm\,2\%$. What is the smallest sample size required to obtain the desired margin of error ?

$$95\%$$
 confidence interval: $p \pm 1.96\sqrt{\frac{p(1-p)}{n}}$

Goal: find the smallest
$$n$$
 such that it guarantees that $1.96\sqrt{\frac{p(1-p)}{n}} \leqslant 0.02$

Conditions For Interval Estimation For The Proportion

- 1. Random Samples
- 2. Normal Condition: the sampling distribution of p needs to be normal
 - $np \ge 10$
 - $nq \geqslant 10$

3. Independence (n < 10% population size)

What to do when the normal condition is not met?

- Wilson score interval
- Jeffreys interval
- Agresti-Coull interval
- Arcsine transformation
- Clopper–Pearson interval (the exact method)