Lecture 2: Sampling Distributions

BMI 713 October 24, 2017 Peter J Park

Estimation of parameters

- So far, we have assumed that the values of the parameters of a probability distributions are known.
- In the real world, these parameters are generally unknown.
- Let's use the observations in a sample to estimate a population parameter.
- **Point estimation** calculates a single number to estimate the population parameter, e.g., ρ for a binomial distribution, μ for a normal distribution.
- Interval estimation specifies a range of values for a parameter

Estimating the population mean

- We would like to estimate the mean height for graduate students at HMS
- How do we estimate the population mean μ ?
- \bullet The obvious approach would be to use the mean of the sample \bar{x} to estimate the unknown population mean μ
- ullet $ar{x}$ is called an estimator of the parameter μ
- For an unbiased estimate, the sample we have the properties of a random sample: i.i.d. - independent and identically distributed.
- The sample that you have is not the only that could have been selected—it is one of many possible samples

- A second sample of n observations could be chosen and its sample mean calculated.
- ullet $ar{x}_1$ and $ar{x}_2$ are not likely to be equal (sample variability)
- Before we use \bar{x} as an estimator of μ , we need to understand its properties.
- If we were to continue selecting samples of size n indefinitely and computing their means, we would end up with a set of values consisting entirely of samples means.
- The sample mean \bar{x} is a random variable, with outcomes $\bar{x}_1, \bar{x}_2, \bar{x}_3, \ldots$
- How does \bar{x} behave?

Sampling distribution

- The probability distribution of all possible sample means is the sampling distribution of the mean.
- Understanding the properties of a theoretical sampling distribution of means makes it possible to draw conclusions based on a single such sample
- It can be shown that the average of the sample means based on repeated samples of size n approaches the population mean μ as the number of samples selected gets large

$$E(\bar{x}) = \mu$$

- ullet We would expect the sample means to cluster around μ
- We would also expect that the larger the sample size n, the more reliable the estimator \bar{x}

$$E\left[\frac{1}{n}\sum_{i=1}^{n}x_{i}\right] = \frac{1}{n}E\left[\sum_{i=1}^{n}x_{i}\right] \quad Var\left[\frac{1}{n}\sum_{i=1}^{n}x_{i}\right] = \frac{1}{n^{2}}Var\left[\sum_{i=1}^{n}x_{i}\right]$$

$$= \frac{1}{n}\sum_{i=1}^{n}E(x_{i}) \qquad = \frac{1}{n^{2}}\sum_{i=1}^{n}Var(x_{i})$$

$$= \frac{1}{n}\sum_{i=1}^{n}\mu \qquad = \frac{1}{n^{2}}\sum_{i=1}^{n}\sigma^{2}$$

$$= \mu \qquad = \frac{\sigma^{2}}{n}$$

- As n gets larger, $Var(\bar{x})$ decreases
- There is less variability among the sample means \bar{x} than there is among the individual observations x

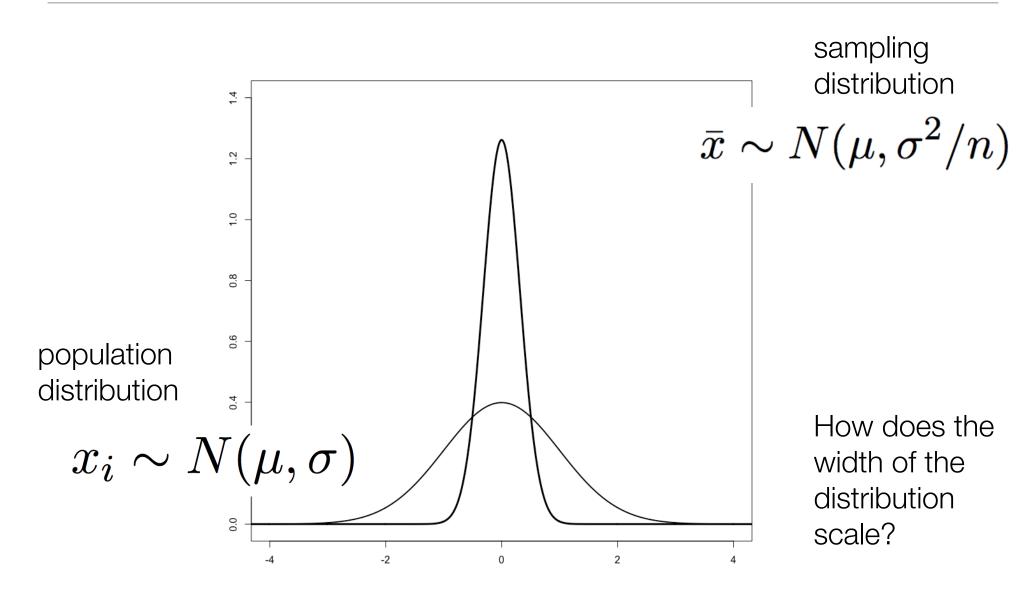
Standard error

- The standard deviation of \bar{x} is σ/\sqrt{n}
- This is called the standard error of the mean

$$\bar{x} \sim N(\mu, \sigma^2/n)$$

- Note that σ^2/n is determined by both the sample size and the degree of variability among the individual observations
- In general, σ quantifies the amount of variability among individuals in a population, while σ/\sqrt{n} quantifies the variability among means of repeated samples drawn from that population

Width of the sampling distribution



The Central Limit Theorem

- If x_i is normally distributed, it can be shown that \bar{x} is also normally distributed
- What if x_i is **not** normally distributed?

$$\bar{x} \sim N(\mu, \sigma^2/n)$$

- Provided that n is large enough, the shape of the sampling distribution is still approximately normal!
- This is called the central limit theorem
- No matter what the underlying distribution of values looks like, inferences about the mean can be based on the normal distribution

What value of *n* is 'large enough' for CLT?

- It depends on the underlying distribution
- If the distribution is itself normal, then n=1 is large enough
- The further the population is from being normally distributed, the larger *n* has to be

Example

- The distribution of serum cholesterol levels for all 20- to 70-year-old males living in the United States has mean μ = 211 mg/100 ml and standard deviation σ = 46 mg/100 ml
- If a sample of size 25 is selected from this population, what is the probability that the sample has a mean of 230 or above?

 What mean value of serum cholesterol level cuts off the lower 10 of the sampling distribution?

Confidence intervals

- ullet In reality, it is not known whether $ar{x}$ is close to μ or not
- How do we measure the variability of the sample mean?
- \bullet Intuitively, if the variability of \bar{x} is large, it is possible that the sample mean could be far from μ
- Recall

$$\bar{x} \sim N(\mu, \sigma^2/n)$$

If n is large,

$$z = \frac{x - \mu}{\sigma / \sqrt{n}}$$

is a standard normal r.v.

For the standard normal distribution,

$$P(-1.96 \le z \le 1.96) = 0.95$$

• Substituting $(\bar{x} - \mu)/(\sigma/\sqrt{n})$ for z

$$P\left(-1.96 \le \frac{\bar{x} - \mu}{\sigma/\sqrt{n}} \le 1.96\right) = 0.95$$

$$P\left(-1.96 \frac{\sigma}{\sqrt{n}} \le \bar{x} - \mu \le 1.96 \frac{\sigma}{\sqrt{n}}\right) = 0.95$$

$$P\left(\mu - 1.96 \frac{\sigma}{\sqrt{n}} \le \bar{x} \le \mu + 1.96 \frac{\sigma}{\sqrt{n}}\right) = 0.95$$

So 95% of all sample means will lie in the interval

$$\left(\mu - 1.96 \frac{\sigma}{\sqrt{n}}, \, \mu + 1.96 \frac{\sigma}{\sqrt{n}}\right)$$

- ullet The statement provides information about the behavior of $\bar{\mathcal{X}}$ if the population mean μ is known
- ullet The problem is that μ is not known it is what we are trying to estimate
- Back to $P\left(-1.96 \frac{\sigma}{\sqrt{n}} \le \bar{x} \mu \le 1.96 \frac{\sigma}{\sqrt{n}}\right) = 0.95$ $P\left(\bar{x} 1.96 \frac{\sigma}{\sqrt{n}} \le \mu \le \bar{x} + 1.96 \frac{\sigma}{\sqrt{n}}\right) = 0.95$
- The probability that the true population mean μ will be contained in

$$\left(\bar{x} - 1.96 \frac{\sigma}{\sqrt{n}}, \, \bar{x} + 1.96 \frac{\sigma}{\sqrt{n}}\right)$$
 is 95%