Lecture Overview

- Estimation concepts
- Example point estimates
- Example interval estimates
- Basic sample size calculations
- Excursion: Confidence Intervals for regression coefficients

Parametric Statistical Inference: Estimation

- **Statistical inference** includes all procedures that **draw conclusions** about an unknown population based on **one set** of *n* sample observations.
- Question: Where is uncertainty in inferential statistics coming from?
 Answer: Drawing a random sample from the underlying population. Drawing different samples will lead to different sets of observed sample data and, therefore, to different estimates.

- Statistical inference deals with:
 - [a] *estimation* of a population characteristic, which usually is a parameter of the underlying population distribution, i.e., μ or σ^2 of the normal distribution, and *quantifying the uncertainty* of the estimate.
 - [b] *testing hypotheses* (initial assumptions) with regards to parameters of an underlying population distribution.
 - Based on the set of sample observations a specific hypothesis can is **rejected** at a given **error probability** α under the assumption that the hypothesis is true (see BBR Chapter 8).
- In both approaches a sample $\{X_1, X_2, ..., X_n\}$ of given size n is drawn and a **sampling statistics** T is selected that is related to the unknown parameter θ of the **population distribution**:

$$\hat{\theta} = T(x_1, x_2, \dots, x_n).$$

The **hat** on top of $\hat{\theta}$ denotes the **estimate** of the unknown population parameter θ .

- \circ For the estimation approach, $\hat{\theta}$ allows expressing that the parameter θ of the population distribution falls with a given *level of certainty* into an interval around θ .
- For the hypothesis testing approach, $\hat{\theta}$ allow us to make statements about the probability that the observed *sampling statistic differs significantly* from an underlying

hypothetical reference population parameter θ_0 (the <u>naught</u> subscript in θ_0 refers to the **assumed** population characteristic).

Statistical Estimation

- <u>Def. Point Estimation:</u> A single number is calculated from the sample and it is used as the **best** estimate $\hat{\theta}$ of some unknown population parameter θ .
- <u>Def. Interval Estimation</u>: In interval estimation the sample is used to identify a *range* $[\hat{\theta}_L, \hat{\theta}_U]$ of estimation bounds for the *unknown population parameter* θ , within which it is believed to be embedded with a *given probability* 1α .

Concept: Point Estimation

- Def. Statistical Estimator and Statistical Estimate:
 - O A statistical estimator is a generic function T of the n random variables $X_1, X_2, ..., X_n$ from a sample. An estimator T is, therefore, also a **random variable**.
 - Once the sample is taken, the observed values of the random variables are known. These are denoted by $x_1, x_2, ..., x_n$.

- The value of the estimator $\hat{\theta} = T(X_1 = x_1, X_2 = x_2, ..., X_n = x_n)$ is known as the **statistical estimate** of the population parameter θ .
- There are *many possible estimation rules* (functional specifications of the estimator) to estimate a population parameter.

For instance, instead of the mean \bar{X} we may use the mode or median to estimate the parameter μ of an underlying population.

- Two criteria to evaluate whether a selected estimation procedure is the best, are
 - [a] the *bias* and
 - [b] the *efficiency*.

Both criteria are based on the deviation of the estimator from its true population value, i.e., $\hat{\theta} - \theta$.

• The *mean estimation error* (the expected variation around the true population value) is:

$$E(\hat{\theta} - \theta) = \sum_{i=1}^{p} (\hat{\theta}_i - \theta) \cdot \Pr(\hat{\theta}_i) \text{ or } E(\hat{\theta} - \theta) = \int_{-\infty}^{\infty} (\hat{\theta}_i - \theta) \cdot f(\hat{\theta}) \cdot d\hat{\theta}.$$

Main challenge is to develop the distribution $\Pr(\hat{\theta}_i)$ or density $f(\hat{\theta})$ of an estimation rule $\hat{\theta} = T(X_1 = x_1, X_2 = x_2, ..., X_n = x_n)$.

- <u>Def. Unbiased Estimator</u>: An estimator $\hat{\theta}$ of a population parameter is said to be unbiased if its expected value is equal to the population parameter. That is, $\hat{\theta}$ is unbiased of $E(\hat{\theta}) = \theta$.
- Example: Biased versus unbiased estimator:

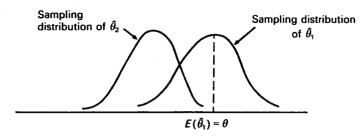


FIGURE 8-3. Sampling distributions for a biased and an unbiased estimator of θ .

A biased estimator may become asymptotically unbiased as the sample size n increases.
 That is, the bias is consistently shrinking.

An example is the biased variance estimator $\frac{1}{n} \cdot \sum_{i=1}^n (x_i - \bar{x})^2$, because for large n we get $\lim_{n \to \infty} n \approx n-1$.

• In how far an *estimator varies around the population value* from sample to sample is measured by the expected squared differences over all possible samples $E\Big[(\hat{\theta}-\theta)^2\Big]$, that is, $Var(\hat{\theta})$.

- Ultimately, we want to have an *unbiased* estimation rule T that, in addition, has the *smallest* possible variability for sample to sample, i.e., the smallest variance.
- Example: Unbiased estimators with smaller versus larger variance:

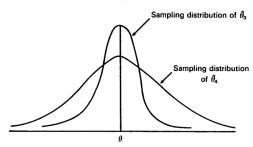


FIGURE 8-4. Sampling distributions for two unbiased estimators of 0.

- <u>Def efficient estimator</u>: An *unbiased* estimation rule T is called *efficient* or best unbiased estimator, if it has the *smallest variance* compared to any other possible *unbiased* estimation rules.
- Which estimation rule T is the most efficient may dependent on the underlying population distribution.
- <u>Def consistency:</u> An estimator is called consistent if for an increasing sample size its estimated value $\hat{\theta}$ approaches the true population value θ and its variance is shrinking.

That is, it *converges in probability* to the true population value:

$$\lim_{n\to\infty} \Pr(|\hat{\theta}_n - \theta| < \delta) = 1$$

for any small, positive value of δ .

• For biased estimation rules with $E(\hat{\theta}) \neq \theta$ the concept of **efficiency does not apply** and their **mean square error** needs to be evaluated

$$MSE = E(\hat{\theta} - \theta)^{2} = \underbrace{\left[E(\hat{\theta}) - \theta\right]^{2}}_{Bias^{2} of T} + \underbrace{E\left[\hat{\theta} - E(\hat{\theta})\right]^{2}}_{Variance of T}$$

• There may be a *tradeoff* between a bias and the variance of an estimator T in the MSE:

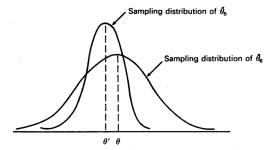


FIGURE 8-5. Difficulties in choosing a potential estimator.

Ultimately, one may *prefer* an estimation rule with a small bias but with a substantially smaller *MSE* compared to alternative unbiased estimation rules.

• To express the variability in original units, one can use the *root mean square error*:

$$RMSE = \sqrt{MSE} = \sqrt{Bias^2 + Variance}$$
.

Note that we *cannot decompose* this expression into the sum

- [a] of the square-root of bias and
- [b] the square-root of the variance, because both terms are jointly under the square root.

Concept: Interval Estimation

- Confidence intervals also depend on the n observed sample observations X₁, X₂,..., X_n.
- These intervals provide more information than a simple point estimators:
 - \circ the *width* defined by a lower and an upper bound of the interval $\left\lceil heta_L, heta_U
 ight
 ceil$
 - o this **width** is associated with the **degree of certainty** of the point estimator $\hat{\theta}$.
- These intervals express the degree of certainty that the true parameter θ is within the **confidence interval**, i.e.,

$$\Pr(\theta \in [\hat{\theta}_L, \hat{\theta}_U]) = 1 - \alpha$$

where lpha is usually small the *error probability* that the *true*

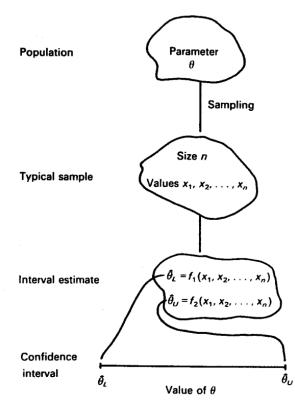


FIGURE 8-6. Interval estimation of a population parameter θ .

population parameter is outside the interval. We want to keep this error probability small.

- The *smaller* the error probability α becomes the *wider* the confidence interval will get, because we are increasing the chance that the interval will be covering the true population parameter.
- See Figure 8-6 conceptionally shows the estimation of the confidence interval:

Examples: Point Estimators

[a] Properties of the Mean Estimation Rule \overline{X}

- For an underlying normal population distribution the arithmetic mean is unbiased and its variance is only 56% of that of the median's variance. It is therefore most efficient.
- Since the mean minimizes the sum of street the squared deviations from the central value, it is highly sensitive to extreme value (leverages).

TABLE 8-1 Point Estimators of μ , π , and σ^2

Population parameters	Point estimator	Formula for point estimate
μ	$ar{X}$	Mean of sample values: $\overline{x} = \frac{1}{n} \sum_{i=1}^{n} x_i$
	Median	Middle value in sample (50%-ile)
	25% Trimmed Mean	Mean of middle 50% of values in sample
	10% Trimmed Mean	Mean of middle 80% of values in sample
π	P	p = x/n where $x =$ number of successes in n trials
σ^2	S^2	$s^2 = \frac{1}{n-1} \sum_{i=1}^{n} (x_i - \bar{x})^2$

 For symmetric distributions with heavy tails the mean may become less efficient than the median.

- For highly *asymmetric distributions* the mean is even less efficient.
- Trimmed means, which disregard the most extreme observations in both tails, are more
 robust because are not affected by these high leverage values.
- The *advantage* of working with the mean is that we know from the central limit theorem its underlying sampling distribution (the normal distribution) as the sample size increases.

[b] Population Proportion

- Assuming the random variables is coded as $X_i = \begin{cases} 1 & \text{for success} \\ 0 & \text{for failure} \end{cases}$.
- Then the population proportion estimation rule is *structurally equivalent* to the arithmetic mean

$$\widehat{\pi} = \frac{\sum_{i=1}^{n} X_i}{n} = \frac{\text{\# of successes}}{\text{\# of trials}}.$$

• Therefore, the properties of the mean apply to the proportion estimator $\hat{\pi}$ for large sample sizes.

[c] Population variance

• The estimator for the sample variance is $S^2 = \frac{1}{n-1} \cdot \sum_{i=1}^{n} (X_i - \bar{X})^2$.

- It is an unbiased estimator (over all potential samples of size n the estimator S^2 will average to the population variance σ^2).
- For a normal distributed underlying population S^2 is the most efficient estimator for σ^2 .
- Division by *n* would lead to a biased estimator, which systematically would *underestimate* the variance. However, for increasing sample sizes it a consistent estimator.
- Review of the reasons:
 - 1. Use of \bar{X} minimizes the sum of the squared deviations.
 - 2. We lose one degree for freedom because, once the mean is known, only n-1 observations need to be available (recall zero sum property $\sum_{i=1}^{n} x_i n \cdot \overline{x} = 0$).

Example: Interval estimation for the population mean

- Review: Standard normal distribution
 - 1. For a **standard normal** distributed variable $(1-\alpha)\times 100\%$ of the observations are within the interval $\left[z_{\alpha/2},z_{\mathrm{l}-\alpha/2}\right]$.

That is,
$$\Pr(z \in [z_{\alpha/2}, z_{1-\alpha/2}]) = 1 - \alpha$$

The quantiles:

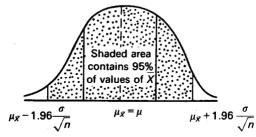


FIGURE 8-7. Sampling distribution of \bar{X} .

 $z_{\alpha/2}$ is in the *left tail* of the standard normal distribution and therefore is *negative*. $z_{1-\alpha/2}$ is in the *right tail* of the standard normal distribution and therefore is *positive*.

- The estimator for the arithmetic mean is distributed as $\bar{X} \sim N(\mu, \sigma/\sqrt{n})$ (recall the central limit theorem).
- Therefore, the confidence interval around the *population expectation* μ is for a *given mean estimation rule*:

$$\begin{split} &\Pr\bigg(z_{\alpha/2} \leq \frac{\overline{X} - \mu}{\sigma/\sqrt{n}} \leq z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\sigma/\sqrt{n} \cdot z_{\alpha/2} \leq \overline{X} - \mu \leq \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(-\overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2} \leq -\mu \leq -\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} - \sigma/\sqrt{n} \cdot z_{\alpha/2} \geq \mu \geq \overline{X} - \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} - \sigma/\sqrt{n} \cdot z_{\alpha/2} \geq \mu \geq \overline{X} - \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2} \geq \mu \geq \overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\bigg) = 1 - \alpha \\ &\Pr\bigg(\overline{X} + \sigma/\sqrt{n} \cdot z_{1-$$

 $\Pr\left(\overline{X} + \sigma/\sqrt{n} \cdot z_{\alpha/2} \le \mu \le \overline{X} + \sigma/\sqrt{n} \cdot z_{1-\alpha/2}\right) = 1 - \alpha$

Parameters $\begin{array}{c}
\chi \\
\hline
\chi \pm 1.96\sigma_{\overline{\chi}}
\end{array}$ $\begin{array}{c}
\chi \pm 1.96\sigma_{\overline{\chi}}
\end{array}$ (Does not encompass μ) $\begin{array}{c}
\chi \pm 1.96\sigma_{\overline{\chi}}
\end{array}$

FIGURE 8-8. Interval estimates constructed from repeated samples of size n.

- Since $z_{0.025} = -1.96$ and $z_{0.975} = 1.96$ at $\alpha = 0.05$ the confidence interval can be visualized by:
- The probability $1-\alpha$ can be interpreted as: $(1-\alpha)\times 100\%$ of the possible samples lead to confidence intervals that **will cover the true but unknown population expectation** μ :

General Rules

- 1. The *smaller the error probability* α (i.e., the confidence level $1-\alpha$ increases) is the *wider the confidence interval* becomes and *vice versa*. Example:
- 2. Furthermore, beside the error probability the **width** of the confidence interval also depends on the **sample size** through the standard error σ/\sqrt{n} of the mean.

As n increases, the confidence interval will shrink. Example:

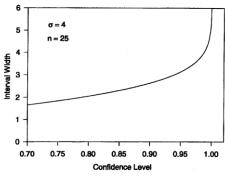


FIGURE 8-10. Effect of confidence level on interval width.

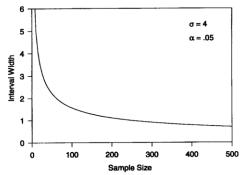


FIGURE 8-11. Effect of sample size on confidence interval width.

Example: Asymptotic Interval for the Expectation from a Non-normal Population

- In this case the underlying sampling distribution of the mean may become difficult to
 evaluate, however, we can make use of the central limit theorem for sufficiently large sample
 size n as then the mean becomes asymptotically normal distributed
- The required sample size depends on the underlying specific circumstances:
 - 1. for *well-behaved population distributions* it may be as low as $n \approx 30$ if just one parameter is estimated from the sample,
 - 2. for the *binomial distribution* it depends on the underlying probability level π . For π not too close to either end of its support $\pi \in [0,1]$, an $n \approx 100$ may be sufficient.

Example: Interval Estimation for the Expectation from a Normal Population with Unknown Standard Deviation

• The unknown population standard deviation σ must be replaced by a sample estimate of the standard deviation $S = \sqrt{\sum_{i=1}^{n} (X_i - \bar{X})^2 / (n-1)}$.

- Since $Z = (\overline{X} \mu) / (\frac{1}{\sqrt{n}} \cdot S)$ contains now **two random variables** (that is, \overline{X} and S) the random variable Z will no longer follow a normal distribution.
 - One can show that Z follows the t-distribution with n-1 degrees for freedom:
 - 1. Like the normal distribution also the t-distribution is **symmetric**.
 - 2. For small degrees of freedom the *t*-distribution has substantially *heavier tails*, that is, it has a *positive kurtosis*.
 - 3. For n > 30, the *t*-distribution can be **approximated** by the normal distribution.
 - \Rightarrow The t-distribution approaches the **standard normal distribution** which does not use degrees of freedom:

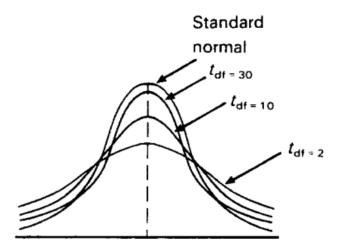


FIGURE 8-12. Normal distribution and the *t* distribution for 2, 10, and 30 df.

• The confidence interval for $n \le 30$ becomes:

$$\Pr\left(\overline{x} + s / \sqrt{n} \cdot t_{\alpha/2, df = n - 1} \le \mu \le \overline{x} + s / \sqrt{n} \cdot t_{1 - \alpha/2, df = n - 1}\right) = 1 - \alpha,$$

where $t_{\alpha/2,df=n-1}$ is the lower quantile and $t_{1-\alpha/2,df=n-1}$ is the upper quantile.

• Note: For n>30 the confidence interval based on the standard normal distribution with $z_{\alpha/2}$ and $z_{1-\alpha/2}$ can be used.

Example: Interval Estimation for the Success Probability of a Binomial Distributed Population

- Because the proportion estimator $\hat{\pi} = \frac{1}{n} \cdot \sum_{i=1}^{n} x_i$ has the structure of an arithmetic mean, we can apply for <u>sufficiently large</u> n the central limit theorem.
- The proportion estimator is unbiased:

$$E(\hat{\Pi}) = \frac{1}{n} \cdot \sum_{i=1}^{n} E(X_i) \text{ with } E(X_i) = 0 \cdot (1 - \pi) + 1 \cdot \pi = \pi$$
$$= \frac{1}{n} \cdot \sum_{i=1}^{n} \pi = \frac{1}{n} \cdot n \cdot \pi = \pi$$

Its variance is:

$$\begin{aligned} Var(\hat{\Pi}) &= Var\Big(\frac{1}{n} \cdot \sum_{i=1}^{n} X_i\Big) \\ &= \frac{1}{n^2} \cdot \sum_{i=1}^{n} \underbrace{Var(X_i)}_{=\pi \cdot (1-\pi)} \\ &= \frac{1}{n^2} \cdot n \cdot \pi \cdot (1-\pi) = \frac{\pi \cdot (1-\pi)}{n} \end{aligned}$$

Since the variance depends on the unknown population probability π , its **estimated value** $\hat{\pi}$ must **substituted** into the equation for the standard error: $s_{\pi} = \sqrt{\frac{\hat{\pi} \cdot (1 - \hat{\pi})}{n}}$

• Using the normal approximation $\widehat{\pi} \sim N\left(\pi, \frac{\widehat{\pi} \cdot (1-\widehat{\pi})}{n}\right)$ the confidence interval for population success probability π becomes

$$\Pr\left(\hat{\pi} + \sqrt{\frac{\hat{\pi} \cdot (1 - \hat{\pi})}{n}} \cdot z_{\alpha/2} \le \pi \le \hat{\pi} + \sqrt{\frac{\hat{\pi} \cdot (1 - \hat{\pi})}{n}} \cdot z_{1 - \alpha/2}\right) = 1 - \alpha$$

Summary of Confidence Intervals

TABLE 8-5 Summary of Point Estimators and Confidence Intervals for π and μ

Population parameter	Point estimator	Formula for confidence intervel	Appropriate conditions
μ	\overline{X}	$\bar{x} \pm z_{\alpha/2}(\sigma/\sqrt{n})$	Exact for any sample size when population standard deviation is known and X normally distributed Approximate when X is not normally distributed but $n > 30$.
μ	\bar{X}	$\overline{x} \pm t_{\alpha/2,n-1f}(s/\sqrt{n})$	Exact when population standard deviation is unknown and X normally distributed Approximate when X is not normal but $n > 30$.
π	P	$p \pm z_{\alpha/2} \sqrt{p(1-p)/n}$	Approximate when $n > 100$.

Sample Size Determination

- Figure 8-11 showed that there is a relationship between the sample size n and the interval width of a confidence interval at a given confidence level $1-\alpha$. These components can be used to determine the required sample size for a given error probability and bounds.
- Steps:

- 1. Determine the estimator for population parameter for which we would like to determine the sample size and its distribution
- 2. Determine the **precision** in terms of the error E, which is half the interval width.
- 3. Determine the confidence level $1-\alpha$.
- Assuming a normal distribution
 - 1. For the population expectation from a normal distribution and a confidence level $1-\alpha$ we get:

The confidence interval bounds are $\bar{x} \pm z_{\alpha/2} \cdot \sigma/\sqrt{n}$; therefore the error E becomes

$$E = |z_{\alpha/2}| \cdot \sigma/\sqrt{n}$$

Solving this expression for sample size leads to $n = \left(\frac{|z_{\alpha/2}| \cdot \sigma}{E}\right)^2$.

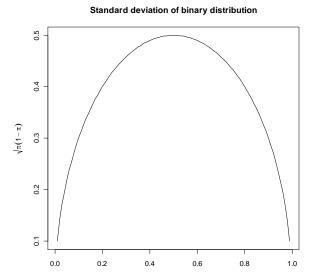
2. To explore the effect of the error on the required sample size let us assume that error becomes twice as large: $E^* = 2 \cdot |z_{\alpha/2}| \cdot \sigma / \sqrt{n}$ we get as required sample size

$$n = \left(\frac{\left|z_{\alpha/2}\right| \cdot \sigma}{2 \cdot E}\right)^2 = \frac{1}{4} \cdot \left(\frac{\left|z_{\alpha/2}\right| \cdot \sigma}{E}\right)^2.$$

Thus just a *quarter* of sample data is needed if we are willing to make the error *twice* as large.

- General rules:
 - 1. As the error E decreases the sample size n increases.
 - 2. As the error probability α *decreases*, the critical tail value $|z_{\alpha/2}|$ becomes larger and, therefore, the required sample size n *increases*.
- Assuming an unknown population variance of normally distributed random variables:
 - 1. This leads to critical values of the t-distribution
 - 2. The *t*-distribution depends on the sample size n. However, for a sample size of n > 30 a normal approximation can be used.
- Assuming a binomial distribution
 - 1. For the proportion estimator $\hat{\pi}$ of a binomial

distribution we get
$$n = \left[\frac{\left| z_{\alpha/2} \right| \cdot \sqrt{\hat{\pi} \cdot (1 - \hat{\pi})}}{E} \right]^2$$



- 2. Since $\hat{\pi}$ is *unknown* before the sample is drawn, it must be determined exogenously (e.g., through past experience).
- 3. Alternatively, the worst case scenario of $\pi=0.5$ can be used, for which the variance $\pi\cdot (1-\pi)$ becomes the largest. This provides a conservative upper bound for n.
- 4. Note, the error E must be substantially smaller than 1/2 in order for the interval covering the support $\pi \in]0,1[$.

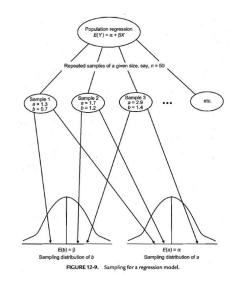
Excursion: Confidence Intervals for Regression Coefficients

• The regression coefficients $b_0, b_1, b_2, ...$ are estimated of the true relationship

$$Y = \beta_0 + \beta_1 \cdot X_1 + \beta_2 \cdot X_2 + \dots + \varepsilon$$

between the dependent variable and Y and the independent variables $X_1, X_2, ...$

- A sampling perspective from an underlying population can be applied:
 - Thus estimated regression coefficients are estimates of the population parameters.



- Each estimated regression coefficient has a sampling distribution with a standard error.
- If the assumptions of regression analysis are satisfied then the estimated regression coefficients are unbiased: $E[b_j] = \beta_j$ for all independent variables X_j .
- The confidence intervals at a given error probability α can be calculated (technical details are covered in "Advanced Data Analysis"):

$$\Pr[\beta_j^{lower,\alpha} \le \beta_j \le \beta_j^{upper,\alpha}] = 1 - \alpha$$

- <u>Lay-person's interpretation (no statistical rigor):</u>
 - If the value 0 is within the confidence interval $0 \in [\beta_j^{lower,\alpha}, \beta_j^{upper,\alpha}]$, then this implies that the true population parameter β_j is not be different from zero.
 - Consequently, the associated independent variable X_j has no influence on the variability of Y.
 - Explore the script StateSchoolConfint.r.