

Mathematical statistics

May 13th, 2019

Review

- Final exam:

Wednesday, 5/29/2019, 10:30am –12:30pm

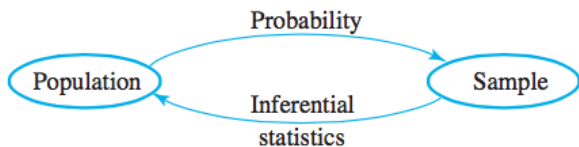
Ewing Hall Room 101

- Course evaluation

Week 1	●	Probability reviews
Week 2	●	Chapter 6: Statistics and Sampling Distributions
Week 4	●	Chapter 7: Point Estimation
Week 7	●	Chapter 8: Confidence Intervals
Week 10	●	Chapter 9, 10: Test of Hypothesis

Chapter 6: Statistics and Sampling Distributions

- 6.1 Statistics and their distributions
- 6.2 The distribution of the sample mean
- 6.3 The distribution of a linear combination



Definition

The random variables X_1, X_2, \dots, X_n are said to form a (simple) random sample of size n if

- 1 the X_i 's are independent random variables
- 2 every X_i has the same probability distribution

Section 6.1: Sampling distributions

- 1 If the distribution and the statistic T is simple, try to construct the pmf of the statistic
- 2 If the probability density function $f_X(x)$ of X 's is known, the
 - try to represent/compute the cumulative distribution (cdf) of T

$$\mathbb{P}[T \leq t]$$

- take the derivative of the function (with respect to t)

Section 6.3: Linear combination of normal random variables

Theorem

Let X_1, X_2, \dots, X_n be independent normal random variables (with possibly different means and/or variances). Then

$$T = a_1X_1 + a_2X_2 + \dots + a_nX_n$$

also follows the normal distribution.

Section 6.3: Computations with normal random variables

If X has a normal distribution with mean μ and standard deviation σ , then

$$Z = \frac{X - \mu}{\sigma}$$

has a standard normal distribution. Thus

$$\begin{aligned}P(a \leq X \leq b) &= P\left(\frac{a - \mu}{\sigma} \leq Z \leq \frac{b - \mu}{\sigma}\right) \\&= \Phi\left(\frac{b - \mu}{\sigma}\right) - \Phi\left(\frac{a - \mu}{\sigma}\right) \\P(X \leq a) &= \Phi\left(\frac{a - \mu}{\sigma}\right) \quad P(X \geq b) = 1 - \Phi\left(\frac{b - \mu}{\sigma}\right)\end{aligned}$$

Section 6.3: Linear combination of random variables

Theorem

Let X_1, X_2, \dots, X_n be independent random variables (with possibly different means and/or variances). Define

$$T = a_1X_1 + a_2X_2 + \dots + a_nX_n,$$

then the mean and the standard deviation of T can be computed by

- $E(T) = a_1E(X_1) + a_2E(X_2) + \dots + a_nE(X_n)$
- $\sigma_T^2 = a_1^2\sigma_{X_1}^2 + a_2^2\sigma_{X_2}^2 + \dots + a_n^2\sigma_{X_n}^2$

Section 6.2: Distribution of the sample mean

Theorem

Let X_1, X_2, \dots, X_n be a random sample from a distribution with mean μ and variance σ^2 . Then, in the limit when $n \rightarrow \infty$, the standardized version of \bar{X} have the standard normal distribution

$$\lim_{n \rightarrow \infty} \mathbb{P} \left(\frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \leq z \right) = \mathbb{P}[Z \leq z] = \Phi(z)$$

Rule of Thumb:

If $n > 30$, the Central Limit Theorem can be used for computation.

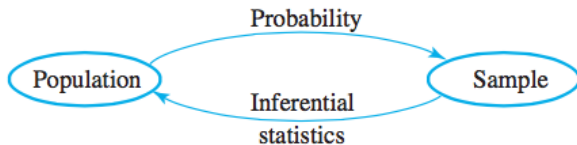
Chapter 7: Point Estimation

7.1 Point estimate

- unbiased estimator
- mean squared error

7.2 Methods of point estimation

- method of moments
- method of maximum likelihood.



Definition

A point estimate $\hat{\theta}$ of a parameter θ is a single number that can be regarded as a sensible value for θ .

$$\begin{array}{ccccc} \text{population parameter} & \implies & \text{sample} & \implies & \text{estimate} \\ \theta & & \implies X_1, X_2, \dots, X_n & \implies & \hat{\theta} \end{array}$$

Definition

The mean squared error of an estimator $\hat{\theta}$ is

$$E[(\hat{\theta} - \theta)^2]$$

Theorem

$$MSE(\hat{\theta}) = E[(\hat{\theta} - \theta)^2] = V(\hat{\theta}) + (E(\hat{\theta}) - \theta)^2$$

Bias-variance decomposition

Mean squared error = variance of estimator + (*bias*)²

Definition

A point estimator $\hat{\theta}$ is said to be an unbiased estimator of θ if

$$E(\hat{\theta}) = \theta$$

for every possible value of θ .

Unbiased estimator

\Leftrightarrow Bias = 0

\Leftrightarrow Mean squared error = variance of estimator

Example

Problem

Consider a random sample X_1, \dots, X_n from the pdf

$$f(x) = \frac{1 + \theta x}{2} \quad -1 \leq x \leq 1$$

Show that $\hat{\theta} = 3\bar{X}$ is an unbiased estimator of θ .

- Let X_1, \dots, X_n be a random sample from a distribution with pmf or pdf

$$f(x; \theta_1, \theta_2, \dots, \theta_m)$$

- Assume that for $k = 1, \dots, m$

$$\frac{X_1^k + X_2^k + \dots + X_n^k}{n} = E(X^k)$$

- Solve the system of equations for $\theta_1, \theta_2, \dots, \theta_m$

Method of moments: example

Problem

Suppose that for a parameter $0 \leq \theta \leq 1$, X is the outcome of the roll of a four-sided tetrahedral die

x	1	2	3	4
$p(x)$	$\frac{3\theta}{4}$	$\frac{\theta}{4}$	$\frac{3(1-\theta)}{4}$	$\frac{(1-\theta)}{4}$

Suppose the die is rolled 10 times with outcomes

4, 1, 2, 3, 1, 2, 3, 4, 2, 3

Use the method of moments to obtain an estimator of θ .

Problem

Let X_1, X_2, \dots, X_n represent a random sample from a distribution with pdf

$$f(x, \theta) = \frac{2x}{\theta + 1} e^{-x^2/(\theta+1)}, \quad x > 0$$

It can be shown that

$$E(X^2 - 1) = \theta$$

Use this fact to construct an estimator of θ based on the method of moments.

Sketch:

$$E(X^2) = \theta + 1$$

MoM:

$$E(X^2) = \frac{X_1^2 + X_2^2 + \dots + X_n^2}{n}$$

Maximum likelihood estimator

- Let X_1, X_2, \dots, X_n have joint pmf or pdf

$$f_{joint}(x_1, x_2, \dots, x_n; \theta)$$

where θ is unknown.

- When x_1, \dots, x_n are the observed sample values and this expression is regarded as a function of θ , it is called the likelihood function.
- The maximum likelihood estimates θ_{ML} are the value for θ that maximize the likelihood function:

$$f_{joint}(x_1, x_2, \dots, x_n; \theta_{ML}) \geq f_{joint}(x_1, x_2, \dots, x_n; \theta) \quad \forall \theta$$

How to find the MLE?

- Step 1: Write down the likelihood function.
- Step 2: Can you find the maximum of this function?
- Step 3: Try taking the logarithm of this function.
- Step 4: Find the maximum of this new function.

To find the maximum of a function of θ :

- compute the derivative of the function with respect to θ
- set this expression of the derivative to 0
- solve the equation

Problem

Let X_1, X_2, \dots, X_n represent a random sample from a distribution with pdf

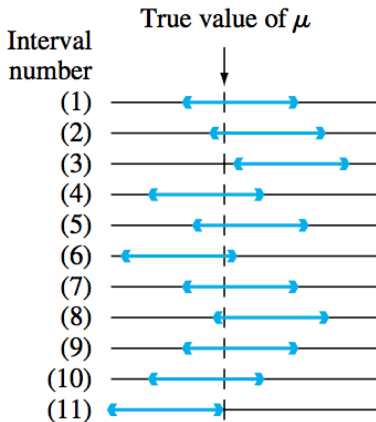
$$f(x, \theta) = \frac{2x}{\theta + 1} e^{-x^2/(\theta+1)}, \quad x > 0$$

Derive the maximum-likelihood estimator for parameter θ based on the following dataset with $n = 10$

17.85, 11.23, 14.43, 19.27, 5.59, 6.36, 9.41, 6.31, 13.78, 11.95

Chapters 8 and 10: Confidence intervals

Interpreting confidence interval



95% confidence interval: If we repeat the experiment many times, the interval contains μ about 95% of the time

Confidence intervals

- By target
 - Chapter 8: Confidence intervals for population means
 - Chapter 8: Prediction intervals for an additional sample
 - Chapter 10: Confidence intervals for difference between two population means
 - independent samples
 - paired samples
- By types
 - (Standard) two-sided confidence intervals
 - One-sided confidence intervals (confidence bounds)
- By distributions of the statistics
 - z-statistic
 - t-statistic

Chapter 8: Confidence intervals

- Section 8.1
 - Normal distribution, σ is known
 - Section 8.2
 - ~~Normal distribution, σ is known~~
 - $n > 40$
 - Section 8.3
 - ~~Normal distribution, σ is known~~
 - n is small
- t -distribution

Assumptions:

- Normal distribution
- σ is known

A **$100(1 - \alpha)\%$ confidence interval** for the mean μ of a normal population when the value of σ is known is given by

$$\left(\bar{x} - z_{\alpha/2} \cdot \frac{\sigma}{\sqrt{n}}, \bar{x} + z_{\alpha/2} \cdot \frac{\sigma}{\sqrt{n}} \right) \quad (8.5)$$

or, equivalently, by $\bar{x} \pm z_{\alpha/2} \cdot \sigma / \sqrt{n}$.

If after observing $X_1 = x_1, X_2 = x_2, \dots, X_n = x_n$ ($n > 40$), we compute the observed sample mean \bar{x} and sample standard deviation s . Then

$$\left(\bar{x} - z_{\alpha/2} \frac{s}{\sqrt{n}}, \bar{x} + z_{\alpha/2} \frac{s}{\sqrt{n}} \right)$$

is a 95% confidence interval of μ

Section 8.3

Let \bar{x} and s be the sample mean and sample standard deviation computed from the results of a random sample from a normal population with mean μ . Then a **100(1 - α)% confidence interval for μ , the one-sample t CI**, is

$$\left(\bar{x} - t_{\alpha/2, n-1} \cdot \frac{s}{\sqrt{n}}, \bar{x} + t_{\alpha/2, n-1} \cdot \frac{s}{\sqrt{n}} \right) \quad (8.15)$$

or, more compactly, $\bar{x} \pm t_{\alpha/2, n-1} \cdot s/\sqrt{n}$.

An **upper confidence bound for μ** is

$$\bar{x} + t_{\alpha, n-1} \cdot \frac{s}{\sqrt{n}}$$

and replacing + by - in this latter expression gives a **lower confidence bound for μ** ; both have confidence level 100(1 - α)%.

- We have available a random sample X_1, X_2, \dots, X_n from a normal population distribution
- We wish to predict the value of X_{n+1} , a single future observation.

A **prediction interval (PI)** for a single observation to be selected from a normal population distribution is

$$\bar{x} \pm t_{\alpha/2, n-1} \cdot s \sqrt{1 + \frac{1}{n}} \quad (8.16)$$

The *prediction level* is $100(1 - \alpha)\%$.

- Independent samples

- 1 X_1, X_2, \dots, X_m is a random sample from a population with mean μ_1 and variance σ_1^2 .
- 2 Y_1, Y_2, \dots, Y_n is a random sample from a population with mean μ_2 and variance σ_2^2 .
- 3 The X and Y samples are independent of each other.

- Paired samples

- 1 There is only one set of n individuals or experimental objects
- 2 Two observations are made on each individual or object

Difference between population means: independent samples

The **two-sample t confidence interval** for $\mu_1 - \mu_2$ with confidence level $100(1 - \alpha)\%$ is then

$$\bar{x} - \bar{y} \pm t_{\alpha/2, v} \sqrt{\frac{s_1^2}{m} + \frac{s_2^2}{n}}$$

A one-sided confidence bound can be calculated as described earlier.

Difference between population means: paired samples

- The paired t CI for μ_D is

$$\bar{d} \pm t_{\alpha/2, n-1} \frac{s_D}{\sqrt{n}}$$

- A one-sided confidence bound results from retaining the relevant sign and replacing $t_{\alpha/2, n-1}$ by $t_{\alpha, n-1}$.

Principles for deriving CIs

If X_1, X_2, \dots, X_n is a random sample from a distribution $f(x, \theta)$, then

- Find a random variable $Y = h(X_1, X_2, \dots, X_n; \theta)$ such that the probability distribution of Y does not depend on θ or on any other unknown parameters.
- Find constants a, b such that

$$P[a < h(X_1, X_2, \dots, X_n; \theta) < b] = 1 - \alpha$$

- Manipulate these inequality to isolate θ

$$P[\ell(X_1, X_2, \dots, X_n) < \theta < u(X_1, X_2, \dots, X_n)] = 1 - \alpha$$

- For μ and X_{n+1}

$$\frac{\bar{X} - \mu}{S/\sqrt{n}} \sim t_{n-1}, \quad \frac{\bar{X} - X_{n+1}}{S\sqrt{1 + 1/n}} \sim t_{n-1}$$

- Difference between two means [independent samples]

$$\frac{(\bar{X} - \bar{Y}) - (\mu_1 - \mu_2)}{\sqrt{\frac{S_1^2}{m} + \frac{S_2^2}{n}}} \sim t_\nu$$

- Difference between two means [paired samples]

$$T = \frac{\bar{D} - \mu_D}{S_D/\sqrt{n}} \sim t_{n-1}$$

Chapters 9 and 10: Tests of hypotheses

Test of hypotheses

- By target
 - Chapter 9: population mean
 - Chapter 10: difference between two population means
 - independent samples
 - paired samples
- By the alternative hypothesis
 - $>$
 - $<$
 - \neq
- By the type of test
 - z-test
 - t-test
- By method of testing
 - Rejection region
 - p-value

In any hypothesis-testing problem, there are two contradictory hypotheses under consideration

- The null hypothesis, denoted by H_0 , is the claim that is initially assumed to be true
- The alternative hypothesis, denoted by H_a , is the assertion that is contradictory to H_0 .

- H_0 will always be stated as an equality claim.
- If θ denotes the parameter of interest, the null hypothesis will have the form

$$H_0 : \theta = \theta_0$$

- θ_0 is a specified number called the *null value*
- The alternative hypothesis will be either:
 - $H_a : \theta > \theta_0$
 - $H_a : \theta < \theta_0$
 - $H_a : \theta \neq \theta_0$

A test procedure is specified by the following:

- A test statistic T : a function of the sample data on which the decision (reject H_0 or do not reject H_0) is to be based
- A rejection region \mathcal{R} : the set of all test statistic values for which H_0 will be rejected
- A type I error consists of rejecting the null hypothesis H_0 when it is true
- A type II error involves not rejecting H_0 when H_0 is false.

Hypothesis testing for one parameter: rejection region method

- 1 Identify the parameter of interest
- 2 Determine the null value and state the null hypothesis
- 3 State the appropriate alternative hypothesis
- 4 Give the formula for the test statistic
- 5 State the rejection region for the selected significance level α
- 6 Compute statistic value from data
- 7 Decide whether H_0 should be rejected and state this conclusion in the problem context

Normal population with known σ

Null hypothesis: $\mu = \mu_0$

Test statistic:

$$Z = \frac{\bar{X} - \mu_0}{\sigma/\sqrt{n}}$$

...

Alternative Hypothesis

$$H_a: \mu > \mu_0$$

$$H_a: \mu < \mu_0$$

$$H_a: \mu \neq \mu_0$$

Rejection Region for Level α Test

$$z \geq z_\alpha \text{ (upper-tailed test)}$$

$$z \leq -z_\alpha \text{ (lower-tailed test)}$$

$$\text{either } z \geq z_{\alpha/2} \text{ or } z \leq -z_{\alpha/2} \text{ (two-tailed test)}$$

Large-sample tests

Null hypothesis: $\mu = \mu_0$

Test statistic:

$$Z = \frac{\bar{X} - \mu_0}{S/\sqrt{n}}$$

...

Alternative Hypothesis

$$H_a: \mu > \mu_0$$

$$H_a: \mu < \mu_0$$

$$H_a: \mu \neq \mu_0$$

Rejection Region for Level α Test

$$z \geq z_\alpha \text{ (upper-tailed test)}$$

$$z \leq -z_\alpha \text{ (lower-tailed test)}$$

$$\text{either } z \geq z_{\alpha/2} \text{ or } z \leq -z_{\alpha/2} \text{ (two-tailed test)}$$

[Does not need the normal assumption]

Null hypothesis: $H_0: \mu = \mu_0$

Test statistic value: $t = \frac{\bar{x} - \mu_0}{s/\sqrt{n}}$

Alternative Hypothesis

$H_a: \mu > \mu_0$

$H_a: \mu < \mu_0$

$H_a: \mu \neq \mu_0$

Rejection Region for a Level α Test

$t \geq t_{\alpha, n-1}$ (upper-tailed)

$t \leq -t_{\alpha, n-1}$ (lower-tailed)

either $t \geq t_{\alpha/2, n-1}$ or $t \leq -t_{\alpha/2, n-1}$ (two-tailed)

[Require normal assumption]

DEFINITION

The ***P*-value** (or *observed significance level*) is the smallest level of significance at which H_0 would be rejected when a specified test procedure is used on a given data set. Once the *P*-value has been determined, the conclusion at any particular level α results from comparing the *P*-value to α :

1. $P\text{-value} \leq \alpha \Rightarrow$ reject H_0 at level α .
2. $P\text{-value} > \alpha \Rightarrow$ do not reject H_0 at level α .

Testing by P-value method

DECISION
RULE BASED
ON THE
P-VALUE

Select a significance level α (as before, the desired type I error probability).
Then reject H_0 if $P\text{-value} \leq \alpha$; do not reject H_0 if $P\text{-value} > \alpha$

Remark: the smaller the P-value, the more evidence there is in the sample data against the null hypothesis and for the alternative hypothesis.

P-values for z-tests

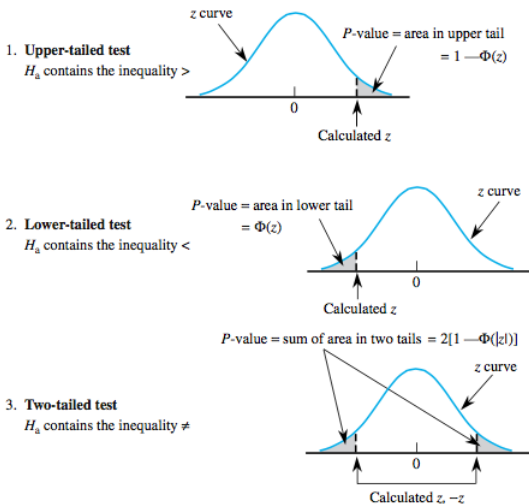


Figure 9.7 Determination of the P -value for a z test

P-values for t -tests

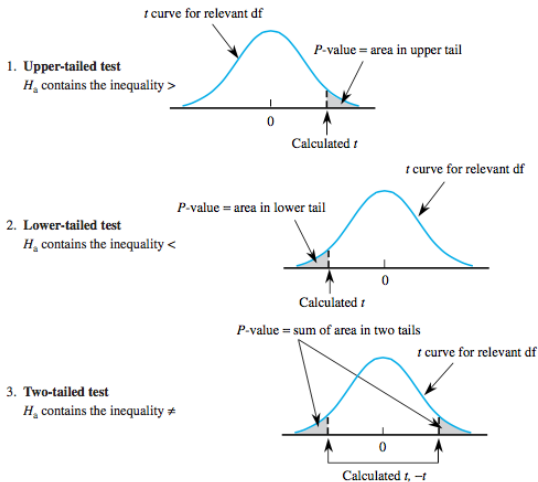


Figure 9.8 P-values for t tests

Testing by rejection region method

- Parameter of interest: $\mu =$ true average activation temperature
- Hypotheses

$$H_0 : \mu = 130$$

$$H_a : \mu \neq 130$$

- Test statistic:

$$z = \frac{\bar{x} - 130}{1.5/\sqrt{n}}$$

- Rejection region: either $z \leq -z_{0.005}$ or $z \geq z_{0.005} = 2.58$
- Substituting $\bar{x} = 131.08$, $n = 25 \rightarrow z = 2.16$.
- Note that $-2.58 < 2.16 < 2.58$. We fail to reject H_0 at significance level 0.01.
- The data does not give strong support to the claim that the true average differs from the design value.

Testing by p-value

1. Parameter of interest: $\mu =$ true average wafer thickness
2. Null hypothesis: $H_0: \mu = 245$
3. Alternative hypothesis: $H_a: \mu \neq 245$
4. Formula for test statistic value: $z = \frac{\bar{x} - 245}{s/\sqrt{n}}$
5. Calculation of test statistic value: $z = \frac{246.18 - 245}{3.60/\sqrt{50}} = 2.32$
6. Determination of P -value: Because the test is two-tailed,
$$P\text{-value} = 2[1 - \Phi(2.32)] = .0204$$
7. Conclusion: Using a significance level of .01, H_0 would not be rejected since $.0204 > .01$. At this significance level, there is insufficient evidence to conclude that true average thickness differs from the target value.

A P-value:

- is not the probability that H_0 is true
- is not the probability of rejecting H_0
- is the probability, calculated assuming that H_0 is true, of obtaining a test statistic value at least as contradictory to the null hypothesis as the value that actually resulted

Testing the difference between two population means

- Setting: independent normal random samples X_1, X_2, \dots, X_m and Y_1, Y_2, \dots, Y_n with known values of σ_1 and σ_2 . Constant Δ_0 .
- Null hypothesis:

$$H_0 : \mu_1 - \mu_2 = \Delta_0$$

- Alternative hypothesis:
 - (a) $H_a : \mu_1 - \mu_2 > \Delta_0$
 - (b) $H_a : \mu_1 - \mu_2 < \Delta_0$
 - (c) $H_a : \mu_1 - \mu_2 \neq \Delta_0$
- When $\Delta = 0$, the test (c) becomes

$$H_0 : \mu_1 = \mu_2$$

$$H_a : \mu_1 \neq \mu_2$$

Difference between 2 means (independent samples)

Proposition

The **two-sample t test** for testing $H_0: \mu_1 - \mu_2 = \Delta_0$ is as follows:

$$\text{Test statistic value: } t = \frac{\bar{x} - \bar{y} - \Delta_0}{\sqrt{\frac{s_1^2}{m} + \frac{s_2^2}{n}}}$$

Alternative Hypothesis Rejection Region for Approximate Level α Test

$$H_a: \mu_1 - \mu_2 > \Delta_0$$

$$t \geq t_{\alpha, v} \text{ (upper-tailed test)}$$

$$H_a: \mu_1 - \mu_2 < \Delta_0$$

$$t \leq -t_{\alpha, v} \text{ (lower-tailed test)}$$

$$H_a: \mu_1 - \mu_2 \neq \Delta_0$$

$$\text{either } t \geq t_{\alpha/2, v} \text{ or } t \leq -t_{\alpha/2, v} \text{ (two-tailed test)}$$

A P -value can be computed as described in Section 9.4 for the one-sample t test.