## Review of Statistics II

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- Estimation
- Properties of Estimators
- Finite and asymptotic properties
- Estimation methods
- Hypothesis testing

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- Inference is based on the sample statistics.
- Two types of estimation: Point vs Interval

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- Two types of estimation: Point vs Interval

- An estimator for a population parameter is simply a function of sample information. Since it is a function of random variables an estimator itself is a random variable.
- The distribution of an estimator is called sampling distribution.
- A certain realization of an estimator based on a sample observation is called an estimate.
- Suppose we want to estimate the mean income of households in Istanbul. For this purpose we can use sample mean which is an estimator for the population mean.
- As we select a new sample it is almost certain that we will obtain a different estimate.
- But on the average we will obtain a value close to the true population mean.

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## Some parameters and estimators

Population parameter	Estimator	Estimate
Mean $(\mu)$	$\overline{X}$	$\bar{x}$
Variance $(\sigma^2)$	$s_X^2$	$s_x^2$
Standard Deviation $(\sigma)$	$s_X$	$s_x$
Success rate $(p)$	$\hat{p}_X$	$\hat{p}_x$

## Properties of Estimators

- How do we decide which estimators to use for inference? We require estimators to have certain properties.
- Finite sample properties: these are valid for any sample size n, unbiasedness and efficiency
- Asymptotic properties: only valid for large samples, e.g., consistency, asymptotic efficiency, asymptotic normality

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 $\theta$ : Unknown population parameter

 $\hat{\theta}$ : a point estimator for  $\theta$ 

Definition: If the mean of the sampling distribution of  $\hat{\theta}$  is equal to the population parameter  $\theta$ , i.e.,

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

then we say that  $\hat{\theta}$  is an unbiased estimator of  $\theta.$ 

#### What does it mean?

- Let us make an hypothetical experiment. Suppose we draw many samples from the population of interest and calculate the estimator for each sample.
- If our estimator is unbiased then the arithmetic mean of our calculated estimates will be equal to the population parameter
- One can show that the following point estimators are unbiased:

$$E(\overline{X}) = \mu, \quad E(s_X^2) = \sigma^2, \quad E(\hat{p}_X) = p$$

 In other words, sample mean is an unbiased estimator of population mean, sample variance is an unbiased estimator of population variance.

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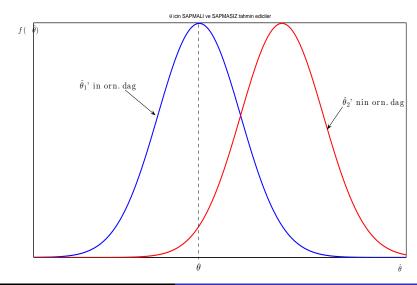
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• If an estimator is not unbiased then it is called biased:

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

- For an unbiased estimator we have  $Bias(\hat{\theta}) = 0$ .
- Unbiasedness does not mean that the value of the estimator is equal to the true parameter value. It just means that on average we would obtain the true parameter.

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### Unbiasedness is not sufficient to obtain good estimators

- In general one can define many unbiased estimators for a parameter.
- We can choose from these unbiased estimators based on their variability, i.e., their variance.
- We would prefer an estimator with smaller variability.
- Definition: let  $\hat{\theta}_1$  and  $\hat{\theta}_2$  be two unbiased estimators for  $\theta$ . If

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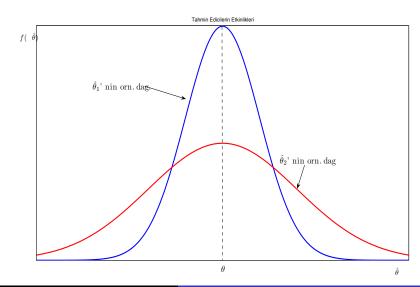
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## Efficiency - Example

- Let  $X_1, X_2, \ldots, X_{10}$  be a random sample from a population with mean  $\mu$  and variance  $\sigma^2$ . Suppose also that there are two estimators for the population mean:  $\hat{\theta}_1 = X_1$  and  $\hat{\theta}_2 = 10^{-1} \sum X_i$ . Are these estimators unbiased? Which one is more efficient?
- Answer: For unbiasedness we must have  $E(\hat{\theta}_1)=\mu$ . Since  $E(\hat{\theta}_1)=E(X_1)=\mu$ ,  $\hat{\theta}_1$  is an unbiased estimator. Similarly,  $E(\hat{\theta}_2)=E(10^{-1}\sum X_i)=\mu$ ,  $\hat{\theta}_2$  is also an unbiased estimator.
- For efficiency we need to calculate their variances.

$$Var(\hat{\theta}_1) = Var(X_1) = \sigma^2 \quad Var(\hat{\theta}_2) = Var(10^{-1} \sum X_i) = 0.01\sigma^2$$

- Obviously  $Var(\hat{\theta}_2) < Var(\hat{\theta}_1)$  thus  $\hat{\theta}_2$  is more efficient than  $\hat{\theta}_1$ .
- Relative efficiency =  $\frac{Var(X_1)}{Var(10^{-1}\sum X_i)} = \frac{\sigma^2}{0.01\sigma^2} = 10$



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- In the previous example our first estimator was defined as the value of the first observation. Note that in this case we would not use the full information contained in the sample.
- As we have shown this estimator is unbiased but inefficient.
   Not matter how large the sample is this variance will not get smaller.
- In most cases we would like our estimators to have good properties as we collect more information, that is as the sample size gets bigger.
- For example, as n becomes large the variance of  $\overline{X}$  gets smaller so that it approaches the true value  $\mu$  at certain rate.
- On the contrary an estimator such as  $X_1$  does not change as n approaches infinity.
- We can eliminate stupid estimators such  $X_1$  by looking at their asymptotic properties.



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- However, it may be relatively easy to obtain the behavior of the estimator as n gets bigger and bigger.
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$$\lim_{n \to \infty} P\left[|\hat{\theta}_n - \theta| < \epsilon\right] = 1 \quad \text{or} \quad \lim_{n \to \infty} P\left[|\hat{\theta}_n - \theta| > \epsilon\right] = 0$$

## then we say that $\hat{\theta}_n$ is a consistent estimator for $\theta$

$$\mathsf{plim}(\hat{\theta}_n) = \theta$$

- This means: as the sample size gets bigger the sampling distribution of  $\hat{\theta}_n$  becomes more concentrated around the true parameter value  $\theta$ . In the limit it becomes degenerate at the true value.
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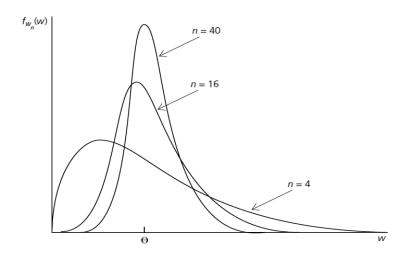
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# Consistency - Sampling Distributions of a Consistent Estimator



$$\operatorname{plim}(\overline{X}_n) = \operatorname{plim}\left(\frac{1}{n}\sum_{i=1}^n X_i\right) = \mu$$

- Since  $Var(\overline{X}_n)$  converges to 0 in the limit sample mean approaches the true value  $\mu$ . Collecting more information leads to smaller and smaller variability.
- In general unbiased estimators with variances converging to zero as n gets bigger are consistent.
- An inconsistent estimator cannot provide information on population parameters.
- An unbiased estimator can be inconsistent and vice versa.



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- Since  $Var(\overline{X}_n)$  converges to 0 in the limit sample mean approaches the true value  $\mu$ . Collecting more information leads to smaller and smaller variability.
- In general unbiased estimators with variances converging to zero as n gets bigger are consistent.
- An inconsistent estimator cannot provide information on population parameters.
- An unbiased estimator can be inconsistent and vice versa.



 The following two conditions are sufficient for an estimator to be consistent:

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$$\lim_{n \to \infty} E(\hat{\theta}_n) = \theta$$

$$2. \quad \lim_{n \to \infty} Var(\hat{\theta}_n) = 0$$

- According to the first condition as the sample size increases expected value becomes closer to the true value. In other words as n goes to infinity bias converges to zero.
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 $\bullet$  Besides the sample mean  $\overline{X}_n$  the following estimators are defined based on a random sample of size n

$$\hat{\mu}_{n}^{1} = \frac{1}{n+1} \sum_{i=1}^{n} X_{t},$$

$$\hat{\mu}_n^2 = \frac{1.02}{n} \sum_{i=1}^n X_t,$$

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$$E(\hat{\mu}_n^1) = E\left(\frac{1}{n+1}\sum_{i=1}^n X_t\right) = \frac{n}{n+1}\mu \Longrightarrow \text{biased}$$

$$Var(\hat{\mu}_{n}^{1}) = Var\left(\frac{1}{n+1}\sum_{i=1}^{n}X_{i}\right) = \frac{n^{2}}{(n+1)^{2}}Var(\overline{X}_{n})$$
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$$n\longrightarrow \infty, \quad \frac{n}{n+1}\mu\longrightarrow \mu, \quad \text{ve} \quad \frac{n}{(n+1)^2}\sigma^2\longrightarrow 0 \Longrightarrow \text{consistent}$$

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Second estimator is biased and inconsistent:

$$E(\hat{\mu}_n^2) = E\left(\frac{1.01}{n}\sum_{i=1}^n X_t\right) = 1.01 E(\overline{X}_n) = 1.01 \mu \Longrightarrow \text{biased}$$
 
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Third estimator is unbiased and inconsistent:

$$\begin{split} E(\hat{\mu}_n^3) &= 0.01 E(X_1) + \frac{0.99}{n-1} E\left(\sum_{i=2}^n X_t\right) = 0.01 \mu + \frac{0.99}{n-1} (n-1) \mu \\ &= \mu \Longrightarrow \text{unbiased} \\ n &\longrightarrow \infty, \quad E\left(0.01 X_1 + \frac{0.99}{n-1} \sum_{i=2}^n X_t\right) \longrightarrow 0.01 X_1 + 0.99 \mu \end{split}$$

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• Recall the definition of probability limit:

$$\mathsf{plim}(Y_n) = \alpha$$

 $\bullet$  The most important property of plim is the following: for any function  $g(Y_n)$  of  $Y_n$ 

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• For example let  $\overline{X}_n > 0$ ,  $g(\overline{X}_n) = \ln(\overline{X}_n)$  .We know that expectation operator does not apply to nonlinear functions:

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- For example let  $Y_1, Y_2, \ldots, Y_n$  be observations on annual income for high school graduates. Let the population mean be  $\mu_Y$  for the high school graduates.
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- This is called *asymptotic normality*.
- Let  $\{Z_n: n=1,2,\ldots,n\}$  be sequence of random variables. Also let  $\Phi(z)$  be cdf of standard normal distribution. If for every z the following condition is satisfied

$$n \longrightarrow \infty \ P(Z \le z) \longrightarrow \Phi(z)$$

then we say that  $Z_n$  has an asymptotic normal distribution.

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- Relies on matching population moments to sample moments
- Suppose we have k population parameters which we want to estimate:  $\theta_1, \theta_1, \dots, \theta_k$
- MoM estimators are found by solving the following system

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### MoM

### Population and sample moments

$\overline{k}$	Population Moments	Sample Moments
1	$\mu_1 = E(X)$	$\hat{\mu}_1 = n^{-1} \sum X_i$
2	$\mu_2 = E(X^2)$	$\hat{\mu}_2 = n^{-1} \sum X_i^2$
3	$\mu_3 = E(X^3)$	$\hat{\mu}_3 = n^{-1} \sum X_i^3$
4	$\mu_4 = E(X^4)$	$\hat{\mu}_4 = n^{-1} \sum X_i^4$
:	:	:
$\underline{k}$	$\mu_k = E(X^k)$	$\hat{\mu}_k = n^{-1} \sum X_i^k$

- Using a random sample of size n drawn from  $N(\mu, \sigma^2)$  find the MoM estimators for  $\mu$  and  $\sigma^2$ .
- Note that there are two unknown parameters. Thus we need two moment conditions.
- Using the first two moments we have

$$E(X) = \mu = \overline{X}$$
  
$$E(X^2) = \mu^2 + \sigma^2 = \frac{1}{n} \sum_{i=1}^n X_i^2$$

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### Maximum Likelihood Estimation - MLE

- Frequently used in practice to estimate fully specified parametric models
- Good asymptotic properties, consistent and asymptotically most efficient
- The distribution of the population (pdf and cdf) has to be known. Strong distributional assumptions may be a weakness.
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- Let  $f(x; \theta)$  represent the distribution (in most cases pdf). Also let  $X_1, X_2, \ldots, X_n$  be a random sample of n observations and let  $x_1, x_2, \ldots, x_n$  be realizations of these random variables.
- Since we have a random sample the joint pdf can be written as follows

$$f(x_1, x_2, \dots, x_n; \theta) = f_1(x_1; \theta) \cdot f_2(x_2; \theta) \cdot, \dots, \cdot f_n(x_n; \theta)$$
$$= \prod_{i=1}^n f(x_i; \theta), \quad i = 1, 2, \dots, n$$

• Now suppose that we observe the random sample but we do not know the population parameters. Likelihood function is simply a function of  $\theta$  given the realization of the random sample  $\mathbf{x}$ :

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- Let  $X_1, X_2, \ldots, X_{10}$  be a random sample drawn from a Bernoulli distribution with success probability p ( $0 ). Also let <math>x_1, x_2, \ldots, x_{10}$  be the observed values of this random sample.
- Let  $y = x_1 + x_2 + \ldots + x_{10}$  denote total number of successes in the random sample. Then the likelihood function is given by:

$$L(p \mid x_1, x_2, \dots, x_{10}) = p^y (1-p)^{n-y}$$

- Suppose that out of 10 trials we observed 6 successes, i.e.,  $y = x_1 + x_2 + \ldots + x_{10} = 6$ .
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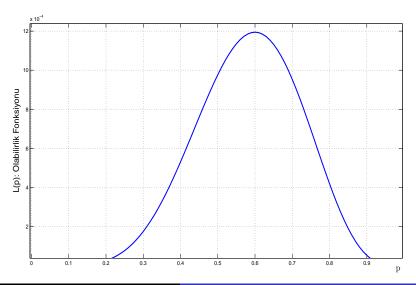


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The likelihood values for the interval  $0.1, 0.2, \dots, 0.9$  are given in the following table:

p	$L(p \mid \mathbf{x}) = p^6 (1 - p)^4$
0.1	0.00000066
0.2	0.00002621
0.3	0.00017503
0.4	0.00053084
0.5	0.00097656
0.6	0.00119439
0.7	0.00095296
8.0	0.00041943
0.9	0.00005314

## MLE - Likelihood Function



- Suppose that  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  is random sample drawn from a Poisson distribution. Find the MLE of  $\lambda$ .
- The probability distribution of  $X \sim Poisson(\lambda)$  is given by:

$$f(x_i; \lambda) = \frac{e^{-\lambda} \lambda^{x_i}}{x_i!}, \quad x_i = 1, 2, 3, \dots, \quad i = 1, 2, \dots, n$$

The loglikelihood function is

$$\ln L(\lambda \mid \mathbf{x}) = \ln \left[ \prod_{i=1}^{n} \frac{e^{-\lambda} \lambda^{x_i}}{x_i!} \right] = \ln \left[ e^{-n\lambda} \prod_{i=1}^{n} \frac{\lambda^{x_i}}{x_i!} \right]$$
$$= -n\lambda + \ln(\lambda) \sum_{i=1}^{n} x_i - \ln \left[ \prod_{i=1}^{n} x_i! \right]$$
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- Suppose that  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  is random sample drawn from a Poisson distribution. Find the MLE of  $\lambda$ .
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• FOC w.r.t.  $\lambda$ :

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• SOC:

$$\frac{\partial^2}{\partial \lambda^2} \ln L(\lambda \mid \mathbf{x}) = -\frac{n\bar{x}}{\lambda^2} < 0, \quad \text{for each } \lambda$$

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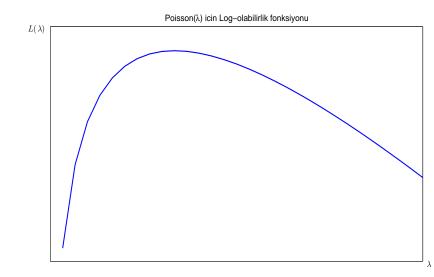
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- Suppose that  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  represent the observed values of a random sample drawn from a normal distribution with mean  $\mu$  and variance  $\sigma^2$ . Find the MLE of  $\mu$  and  $\sigma^2$ .
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#### MLE - Another Example

- Suppose that  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  represent the observed values of a random sample drawn from a normal distribution with mean  $\mu$  and variance  $\sigma^2$ . Find the MLE of  $\mu$  and  $\sigma^2$ .
- $X \sim N(\mu, \sigma^2)$  the pdf is:

$$f(x_i; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2}(x_i - \mu)^2\right), \quad -\infty < x_i < \infty,$$

 We first need to find the joint pdf and the loglikelihood function using the independence property.



Likelihood function is

$$L(\mu, \sigma^{2} \mid \mathbf{x}) = \prod_{i=1}^{n} f(x_{i}; \mu, \sigma^{2}) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma^{2}}} \exp\left(-\frac{1}{2\sigma^{2}}(x_{i} - \mu)^{2}\right)$$

$$= \left(\frac{1}{\sqrt{2\pi\sigma^{2}}}\right)^{n} \prod_{i=1}^{n} \exp\left(-\frac{1}{2\sigma^{2}}(x_{i} - \mu)^{2}\right)$$

$$= (2\pi\sigma^{2})^{-n/2} \exp\left(-\frac{1}{2\sigma^{2}}\sum_{i=1}^{n}(x_{i} - \mu)^{2}\right)$$

Loglikelihood function is

$$\ln L(\mu, \sigma^2 \mid \mathbf{x}) = -\frac{n}{2} \ln(2\pi) - \frac{n}{2} \ln(\sigma^2) - \frac{1}{2\sigma^2} \sum_{i=1}^{n} (x_i - \mu)^2$$



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• FOC:

$$\frac{\partial}{\partial \mu} \ln L(\mu, \sigma^2 \mid \mathbf{x}) = \frac{1}{\sigma^2} \sum_{i=1}^n (x_i - \mu) = 0$$

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By soling this system we obtain

$$\hat{\mu}_{mle} = \overline{X}, \quad \hat{\sigma}_{mle}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \overline{X})^2$$

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## Example continued

#### SOC

$$\frac{\partial^2}{\partial \mu^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) = -\frac{n}{\sigma^2}$$

$$\frac{\partial^2}{\partial \mu \partial \sigma^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) = -\frac{1}{\sigma^4} \sum_{i=1}^n (x_i - \mu)$$

$$\frac{\partial^2}{\partial (\sigma^2)^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) = \frac{n}{2\sigma^4} - \frac{1}{\sigma^6} \sum_{i=1}^n (x_i - \mu)^2$$

$$\frac{\partial^2}{\partial \sigma^2 \partial \mu} \ln L(\mu, \sigma^2 \mid \mathbf{x}) = -\frac{1}{\sigma^4} \sum_{i=1}^n (x_i - \mu)$$

### Example continued

Evaluating the Hessian at the MLE we obtain

$$H|_{\hat{\mu}_{mle}, \hat{\sigma}_{mle}^2} = \begin{bmatrix} \frac{\partial^2}{\partial \mu^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) & \frac{\partial^2}{\partial \mu \partial \sigma^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) \\ \frac{\partial^2}{\partial \sigma^2 \partial \mu} \ln L(\mu, \sigma^2 \mid \mathbf{x}) & \frac{\partial^2}{\partial (\sigma^2)^2} \ln L(\mu, \sigma^2 \mid \mathbf{x}) \end{bmatrix}$$
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Since this matrix is negative definite SOC will be satisfied.

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- Invariance: Let  $\hat{\theta}_{mle}$  be the MLE of  $\theta$ . Also let  $\gamma = g(\theta)$  be a function of  $\theta$  then the MLE of  $\gamma$  is  $\hat{\gamma}_{mle} = g(\hat{\theta}_{mle})$ .
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where

$$\sigma_{\theta}^2 = \frac{1}{I(\theta)}, \quad I(\theta) = E_{\theta} \left[ \frac{\partial}{\partial \theta} \ln L(\theta \mid \mathbf{x}) \right]^2$$

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#### Definitions

- ullet Let the unknown population parameter be heta
- Null Hypothesis: this is the hypothesis that we believe to be true until evidence suggests the otherwise. Contradiction of the alternative hypothesis. Denoted by the symbol  ${\cal H}_0$
- Alternative Hypothesis: is the hypothesis we wish to support. If the null hypothesis is false, the alternative hypothesis must be true. Denoted by the symbol  $H_1$  or  $H_a$
- ullet  $H_0$  specifies hypothesized values for one or more population parameters.
- For example, we might wish to test the null hypothesis that a population mean is equal to 10, hoping to show, in fact that the mean exceeds 10.
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- Two decisions: Fail to reject ("accept") the null hypothesis or reject the null hypothesis in favor of the alternative.
- We need to develop a decision rule using the sample.
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- ullet The probability of Type I Error is denoted lpha and called the significance level.
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DECISION		
	$H_0$ TRUE	$H_0$ FALSE
ACCEPT $H_0$	Correct Decision	Type II Error
	$probability \! = 1 - \alpha$	$probability {= \beta}$
REJECT $H_0$	Type I Error probability= $\alpha$	$\begin{array}{c} \text{Correct Decision} \\ \text{probability} = 1 - \beta \end{array}$

```
\alpha=P(\text{reject }H_0\mid H_0\text{ is true})\text{: Significance Level} \beta=P(\text{fail to reject }H_0\mid H_0\text{ is false})\text{: probability of Type II Error} 1-\beta=P(\text{reject }H_0\mid H_0\text{ is false})\text{: Power of test}
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- $\alpha \downarrow \beta \uparrow (1 \beta) \downarrow$
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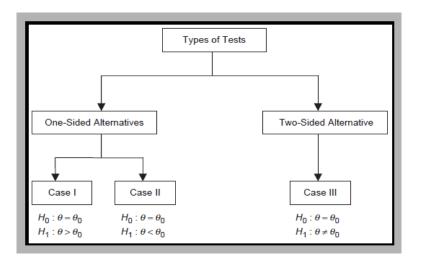
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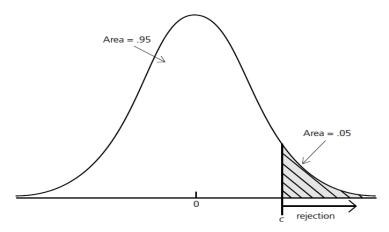
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# Types of tests



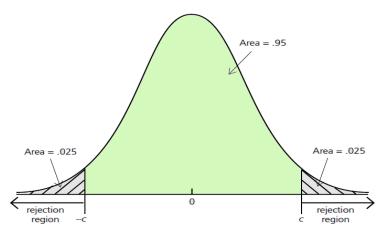
# One-sided Alternative - Right Tail Test

Rejection region for a 5% significance level test against the one-sided alternative  $\mu > \mu_0$ .



#### Two-sided Alternative

Rejection region for a 5% significance level test against the two-sided alternative  $H_1$ :  $\mu \neq \mu_0$ .



- Test results may depend on the significance level chosen by the researcher.
- Instead of choosing a significance level prior to the testing we may ask the following question: What is the largest significance level at which we could carry out the test and still fail to reject the null hypothesis?
- This is called the p-value
- ullet A small p-value indicates strong evidence against  $H_0$
- If  $p-value < \alpha$   $H_0$  can be rejected at significance level  $\alpha$
- For example, suppose that a large random sample (n>120) is drawn from a  $X\sim N(\mu,\sigma^2)$  population and we want to test  $H_0:\mu=0$  against one-sided alternative  $H_1:\mu>0$

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Our test statistic is

$$T = \frac{\overline{X}}{s/\sqrt{n}}$$

- Suppose that the value of T for out sample is t=1.52
- $\bullet$  What is the p-value? i.e., the largest significance level at which we would fail to reject  $H_0$
- $\bullet$  ... or the smallest significance level at which we would reject  ${\cal H}_0$
- Since our sample is large enough so that the distribution of the test statistic is standard normal we can write

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Our test statistic is

$$T = \frac{\overline{X}}{s/\sqrt{n}}$$

- Suppose that the value of T for out sample is t=1.52
- ullet What is the p-value? i.e., the largest significance level at which we would fail to reject  $H_0$
- ullet ... or the smallest significance level at which we would reject  $H_0$
- Since our sample is large enough so that the distribution of the test statistic is standard normal we can write

$$p - value = P(T > 1.52 \mid H_0) = 1 - \Phi(1.52) = 0.065$$



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The p-value when t = 1.52 for the one-sided alternative  $\mu > \mu_0$ .

