

# Point Estimators

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# Three famous frequentist approaches for point estimation

- **Method of Moments**
- Least Squares Estimation
- Maximum Likelihood Estimation

# Method of Moments

- The method of moments starts by equating the sample moments (e.g., mean, variance) with their corresponding population moments.
- We derive equations by setting the sample moments equal to the corresponding theoretical moments.
- The resulting system of equations can be solved to estimate the parameters of the distribution.
- The number of equations needed depends on the number of parameters to be estimated.

## Gaussian model: Mean Estimator

- Let's consider a random sample  $X_1, X_2, \dots, X_n$  from a Gaussian distribution with mean  $\mu$  and variance  $\sigma^2$ .
- The first population moment is the mean,  $\mu_1 = E(X) = \mu$ .
- The first sample moment is the sample mean,  $m_1 = \frac{1}{n} \sum_{i=1}^n X_i$ .
- Equating the population and sample moments, we have  $\mu = \frac{1}{n} \sum_{i=1}^n X_i$ , which gives us the mean estimator  $\hat{\mu}_{MM} = \frac{1}{n} \sum_{i=1}^n X_i$ .

# Derivation of Variance Estimator

## Definition of the r-th moment

- if  $X$  is discrete  $E(X^r) = \sum_i p(x_i) x_i^r$
- if  $X$  is continuous in  $A$   $E(X^r) = \int_A f(x) x^r dx$
- The second population moment is  $\mu_2 = E(X^2) = \frac{1}{n} \sum_{i=1}^n X_i^2$ .
- We know that:  $Var(X) = E(X^2) - (E(X))^2$
- Equating the population and sample moments, we have  $\sigma^2 = \frac{1}{n} \sum_{i=1}^n (X_i)^2 - \left( \frac{1}{n} \sum_{i=1}^n (X_i) \right)^2$ .
- Simplifying the equation, we obtain the variance estimator  $\hat{\sigma}_{MM}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X})^2$ .

# Advantages and Limitations

- Advantages of the method of moments include its simplicity, ease of implementation, and intuitive interpretation
- Is fairly simple and yields consistent estimators (under very weak assumptions)
- It may not always yield the most efficient estimators, especially in small samples or complex models
- It often yields biased estimators
- the suitability of the method depends on the specific problem and the available data

# Likelihood Function

- The likelihood function is a function of the parameters of a statistical model, given the observed data.
- It measures the likelihood or plausibility of the observed data for different parameter values.
- For a random sample of independent and identically distributed (i.i.d.) observations, the likelihood function is the product of the probability density function (pdf) or probability mass function (pmf) for each observation.

# Likelihood Function

## Likelihood Function

The likelihood function, denoted as  $\mathcal{L}(\theta)$ , measures the probability of observing the data given the parameter values  $\theta$ .

- For independent and identically distributed (i.i.d.) observations of a statistical model with density function  $f(\cdot; \theta)$ , the likelihood function is the product of the individual densities.
- If  $X_1, X_2, \dots, X_n$  are i.i.d. random variables, the likelihood function is given by:

$$\mathcal{L}(\theta) = \prod_{i=1}^n f(X_i; \theta)$$



# Properties of the Likelihood Function

- The likelihood function possesses several important properties:
  - ▶ **Non-negativity:** The likelihood function is non-negative for all parameter values.
  - ▶ **Monotonicity:** As the parameter values move away from the true parameter values, the likelihood function decreases.
  - ▶ **Scale invariance:** Multiplying the likelihood function by a constant factor does not change the relative likelihoods of different parameter values.
  - ▶ **Likelihood principle:** The likelihood function contains all the information about the unknown parameters that is available in the data.

# Log-Likelihood Function

## Log-Likelihood Function

The log-likelihood function, denoted as  $\ell(\theta)$ , is the natural logarithm of the likelihood function. It is often easier to work with than the likelihood function itself.

- Taking the logarithm helps simplify calculations and does not change the location of the maximum point.
- The log-likelihood function is given by:

$$\ell(\theta) = \sum_{i=1}^n \ln(f(X_i; \theta))$$

# Score Function

- The score function measures the sensitivity of the log-likelihood function with respect to the parameters of interest.
- It provides information about the direction and magnitude of the parameter effects.
- Mathematically, the score function is defined as:

$$S(\theta) = \frac{\partial}{\partial \theta} \ln \mathcal{L}(\theta|\mathbf{x})$$

where  $\mathcal{L}(\theta|\mathbf{x})$  is the likelihood function,  $\theta$  is the parameter of interest, and  $\mathbf{x}$  is the observed data.

# Fisher Information

- The Fisher information quantifies the amount of information provided by the data about the parameters of interest.
- It measures the precision or uncertainty of the parameter estimates.
- The Fisher information matrix is defined as:

$$\mathcal{I}(\theta) = -\mathbb{E} \left[ \frac{\partial^2}{\partial \theta^2} \ln \mathcal{L}(\theta|\mathbf{x}) \right]$$

where  $\mathbb{E}$  denotes the expectation operator.

# Point Estimators

Three famous frequentist approaches:

- Method of Moments
- Least Squares Estimation
- **Maximum Likelihood Estimation**

# The Maximum Likelihood Estimator

- Maximum Likelihood Estimation (MLE) is a method used to estimate the parameters of a statistical model based on observed data.
- It involves finding the parameter values that maximize the likelihood function, which measures the probability of observing the data given the parameter values.
- MLE is widely used in various fields, including statistics, econometrics, and machine learning.

# Maximum Likelihood Estimation

## Maximum Likelihood Estimation (MLE)

Maximum Likelihood Estimation involves finding the parameter values that maximize the likelihood function (or equivalently, the log-likelihood function).

- The MLE estimates, denoted as  $\hat{\theta}_{\text{MLE}}$ , are obtained by solving the equation  $\frac{\partial \ell(\theta)}{\partial \theta} = 0$ .
- In some cases, it may be easier to maximize the log-likelihood function numerically using optimization algorithms.

# Example: MLE for Bernoulli Distribution

## Step 1: Likelihood Function

Consider a Bernoulli distribution with parameter  $p$ . Let  $X_1, X_2, \dots, X_n$  be independent and identically distributed (i.i.d.) random variables with the following probability mass function:

$$P(X_i = x_i) = \begin{cases} p & \text{if } x_i = 1 \\ 1 - p & \text{if } x_i = 0 \end{cases}$$

The likelihood function, denoted by  $\mathcal{L}(p)$ , can be expressed as the joint probability mass function of the observations:

$$\mathcal{L}(p) = P(X_1 = x_1, X_2 = x_2, \dots, X_n = x_n) = p^{\sum_{i=1}^n x_i} (1 - p)^{n - \sum_{i=1}^n x_i}$$



## Example: MLE for Bernoulli Distribution

### Step 2: Log-Likelihood Function

To simplify the calculations, we take the logarithm of the likelihood function to obtain the log-likelihood function, denoted by  $\ell(p)$ :

$$\ell(p) = \log \mathcal{L}(p) = \sum_{i=1}^n x_i \log(p) + (n - \sum_{i=1}^n x_i) \log(1 - p)$$

## Example: MLE for Bernoulli Distribution

### Step 3: Maximizing the Log-Likelihood

$$\frac{d\ell(p)}{dp} = \frac{\sum_{i=1}^n x_i}{p} - \frac{n - \sum_{i=1}^n x_i}{1-p} = 0$$

$$\frac{\sum_{i=1}^n x_i}{p} = \frac{n - \sum_{i=1}^n x_i}{1-p}$$

$$\sum_{i=1}^n x_i - p \sum_{i=1}^n x_i = pn - p \sum_{i=1}^n x_i$$

$$\sum_{i=1}^n x_i = pn$$

$$\hat{p}_{MLE} = \frac{\sum_{i=1}^n x_i}{n}$$

## Example: MLE for Normal Distribution

### Step 1: Likelihood Function

Consider a normal distribution with unknown mean  $\mu$  and variance  $\sigma^2$ . Let  $X_1, X_2, \dots, X_n$  be independent and identically distributed (i.i.d.) random variables following a normal distribution. The likelihood function, denoted by  $\mathcal{L}(\mu, \sigma^2)$ , can be expressed as the joint density function of the observations:

$$\mathcal{L}(\mu, \sigma^2) = \prod_{i=1}^n \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(X_i - \mu)^2}{2\sigma^2}}$$

## Step 2: Log-Likelihood Function

To simplify the calculations, we take the logarithm of the likelihood function to obtain the log-likelihood function, denoted by  $\ell(\mu, \sigma^2)$ :

$$\ell(\mu, \sigma^2) = \sum_{i=1}^n \left( -\frac{1}{2} \log(2\pi\sigma^2) - \frac{(X_i - \mu)^2}{2\sigma^2} \right)$$

## Example: MLE for Normal Distribution

### Step 3: Maximizing the Log-Likelihood

To find the maximum likelihood estimator, we differentiate the log-likelihood function with respect to  $\mu$  and  $\sigma^2$  and set them equal to zero:

$$\frac{\partial \ell(\mu, \sigma^2)}{\partial \mu} = \frac{\sum_{i=1}^n (X_i - \mu)}{\sigma^2} = 0$$

$$\frac{\partial \ell(\mu, \sigma^2)}{\partial \sigma^2} = \frac{\sum_{i=1}^n (X_i - \mu)^2}{2\sigma^4} - \frac{n}{2\sigma^2} = 0$$

## Step 4: Solving for Maximum Likelihood Estimators

Solving the equations, we obtain the maximum likelihood estimators:

$$\hat{\mu}_{MLE} = \frac{1}{n} \sum_{i=1}^n X_i$$

$$\hat{\sigma}_{MLE}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \hat{\mu}_{MLE})^2$$

# Asymptotic properties of MLE

- The MLE possesses several desirable properties:
  - ▶ Consistency: The MLE converges to the true parameter value as the sample size increases.
  - ▶ Asymptotic Normality: The MLE is asymptotically normally distributed.
  - ▶ Efficiency: The MLE achieves the Cramér-Rao lower bound for the variance of an unbiased estimator.

# Consistency of the MLE

- Let  $\hat{\theta}_{\text{MLE}}$  be the Maximum Likelihood Estimator for parameter  $\theta$ .
- We say that  $\hat{\theta}_{\text{MLE}}$  is consistent if:

$$\lim_{n \rightarrow \infty} \Pr(|\hat{\theta}_{\text{MLE}} - \theta| > \epsilon) = 0 \quad \text{for all } \epsilon > 0.$$

- In simpler terms, the probability that  $\hat{\theta}_{\text{MLE}}$  deviates from  $\theta$  by more than  $\epsilon$  approaches 0 as the sample size  $n$  increases.



# Asymptotic normality of the MLE

- Let  $\hat{\theta}_{MLE}$  be the Maximum Likelihood Estimator for the parameter  $\theta$ .
- It can be proved that  $\hat{\theta}_{MLE}$  is asymptotically normal. Namely, as the sample size  $n$  approaches infinity:

$$\sqrt{n}(\hat{\theta}_{MLE} - \theta) \xrightarrow{d} \mathcal{N}(\theta, \mathcal{I}(\theta)^{-1})$$

where  $\xrightarrow{d}$  denotes convergence in distribution and  $\mathcal{I}(\theta)$  is the Fisher Information matrix.

Note that we can calculate the standard error of the MLE as  $\mathcal{I}(\theta)^{-1}$

## Example: Asymptotic normality of the Bernulli MLE

The log-likelihood function for a sample of  $n$  independent and identically distributed (i.i.d.) observations  $X_1, X_2, \dots, X_n$  from the Bernoulli distribution is:

$$\ell(p) = \log \prod_{i=1}^n f(X_i; p) = \sum_{i=1}^n \log f(X_i; p)$$

Taking the derivative of the log-likelihood function with respect to  $p$ , we get:

$$\frac{d\ell(p)}{dp} = \frac{1}{p} \sum_{i=1}^n X_i - \frac{1}{1-p} \sum_{i=1}^n (1 - X_i)$$

Taking the derivative of the score function we get:

$$\frac{d^2\ell(p)}{dp^2} = -\frac{1}{p^2} \sum_{i=1}^n X_i - \frac{1}{(1-p)^2} \sum_{i=1}^n (1 - X_i)$$

Since  $X_i$  follows a Bernoulli distribution with parameter  $p$ , we have:

$$\mathbb{E}[X_i] = p \quad \text{and} \quad \mathbb{E}[1 - X_i] = 1 - p$$

Substituting these values, we get:

$$\mathbb{E} \left[ \frac{d^2 \ell(p)}{dp^2} \right] = -\frac{1}{p^2} \sum_{i=1}^n p - \frac{1}{(1-p)^2} \sum_{i=1}^n (1-p)$$

Simplifying, we have:

$$\mathbb{E} \left[ \frac{d^2 \ell(p)}{dp^2} \right] = -\frac{n}{p} - \frac{n}{1-p}$$

Finally, the Fisher Information Matrix is the negative expected value of the second derivative of the log-likelihood function:

$$\mathcal{I}(p) = -\mathbb{E} \left[ \frac{d^2 \ell(p)}{dp^2} \right] = \frac{n}{p(1-p)}.$$

Therefore,

$$\hat{p}_{MLE} \xrightarrow{d} \mathcal{N} \left( p, \frac{p(1-p)}{n} \right)$$

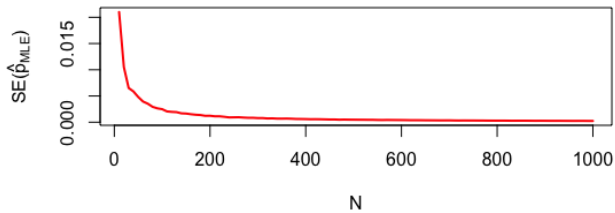
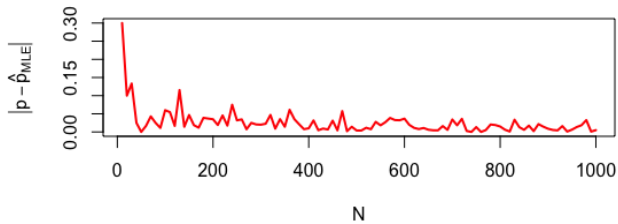
## Consistency: Proof in R

```
library(latex2exp)
N <- seq(10,1000,length.out=100)
p0 <- 0.4

# Generate many random samples of increasing size
and compute MLE.
samples <- lapply(1:100,function(j) rbinom(N[j],
size = 1, prob = p0) )

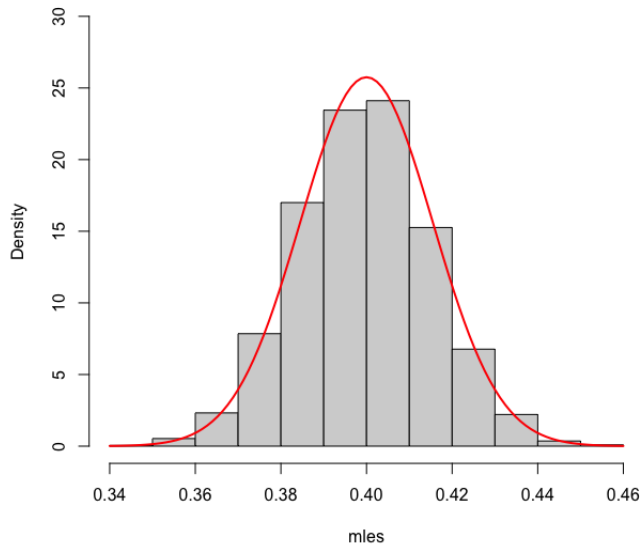
mles<-sapply(1:100, function(j)
mean(unlist(samples[j])))
# Compute the standard error.
se<-(mles*(1-mles))*(N^(-1))

# Plot the standard error values of the MLEs as
the sample size increases.
plot(N,abs(mles-p0), type = "l", lwd=2, col="red",
xlab="N", ylab = TeX("$|p-\hat{p}_{MLE}|$"))
plot(N,se, type = "l", lwd=2, col="red",xlab="N",
ylab = TeX("$SE(\hat{p}_{MLE})$"))
```



## Asymptotic Normality: Proof in R

```
# Plot the asymptotically normal distribution.  
N <- 1000  
p0 <- 0.4  
  
# Generate many random samples of size N and  
compute MLE.  
mles <- replicate(10000, mean(rbinom(N, size = 1,  
prob = p0)))  
  
# Plot histogram of MLEs.  
hist(mles, freq = FALSE, ylim=c(0,30))  
curve(dnorm(x, mean = p0, sd = sqrt((p0 * (1 - p0)) / N)),  
      add=TRUE, col="red", lwd=2)
```



# Efficiency of the MLE

- The CRLB gives a lower bound on the variance of any unbiased estimator.
- By asymptotic Normality of the MLE:

$$\sqrt{n}(\hat{\theta}_{MLE} - \theta) \xrightarrow{d} \mathcal{N}(\theta, \mathcal{I}(\theta)^{-1})$$

we conclude that the variance of the MLE is equal to the CRLB, proving its efficiency.



# Summary on the MLE

- Maximum Likelihood Estimation is a powerful method for estimating parameters in statistical models.
- It involves maximizing the likelihood function (or log-likelihood function) to obtain parameter estimates.
- MLE estimators possess desirable properties, such as consistency and asymptotic normality.
- These properties make the MLE a reliable and powerful tool for parameter estimation.