Lecture 17 Maximum Likelihood Estimation (MLE)

BIO210 Biostatistics

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Intuition over MLE

Experiment: A coin, with an unknown $\mathbb{P}(H) = p$, was flipped 10 times. The outcome is HHHTHHHTHH.

Question: What is your best guess for p?

Thinking: Given the data/observation we have, what values should p take such that our data/observation is most likely to occur ?

Aim: find the value that maximise our chance of observing the data, and use that value as our best guess/estimate for p.

$$\mathcal{L}: \mathbb{P} \left(\mathsf{obs.} \mid \mathbb{P} \left(H \right) = p \right)$$

Estimators of Parameters

• Parameter space Ω : the set of all possible values of a parameter θ or of a vector of parameters $(\theta_1, \theta_2, \theta_3, ..., \theta_k)$ is called the parameter space.

- Bernoulli: $\theta = p$, $\Omega = \{p \mid 0 \leqslant p \leqslant 1\}$
- Binomial: $\theta_1=n, \theta_2=p, \ \Omega=\{(n,p) \mid n=2,3,..., \text{a finite number}; 0\leqslant p\leqslant 1\}$
- Poisson: $\theta = \lambda$, $\Omega = \{\lambda \mid \lambda \geqslant 0\}$
- Normal (Gaussian): $\theta_1 = \mu, \theta_2 = \sigma^2, \ \Omega = \{(\mu, \sigma^2) \mid \mu \in \mathbb{R}, \sigma^2 \geqslant 0\}$
- We refer to an estimator of a parameter θ as $\hat{\boldsymbol{\theta}}$. An estimator $\hat{\boldsymbol{\theta}}$ of a parameter θ is unbiased if $\mathbb{E}\left[\hat{\boldsymbol{\theta}}\right]=\theta$. For example, $\hat{\boldsymbol{\mu}}=\bar{X}$ is an unbiased estimator for μ .

Maximum Likelihood Estimation (MLE)

- Maximum likelihood estimation (MLE) is a technique used for estimating the parameters of a given distribution, using some observed data.
- Introduced by R.A. Fisher in 1912.
- MLE can be used to estimate parameters using a limited sample of the population, by finding particular values so that the observation is the most likely result to have occurred.

Maximum Likelihood Estimation (MLE)

Formal definition

Let $x_1, x_2, x_3, ..., x_n$ be observations from n **i.i.d** random variables $(X_1, X_2, X_3, ..., X_n)$ drawn from a probability distribution f_0 , where f_0 is known to be from a family of distributions f that depend on some paramters θ . For example, f_0 could be known to be from the family of normal distributions f, which depend on parameters μ and σ^2 , and $x_1, x_2, x_3, ..., x_n$ would be observations from f_0 . The goal of MLE is to maximise the likelihood function:

$$\mathcal{L}(\theta; x_1, x_2, x_3, ..., x_n) = f(x_1, x_2, x_3, ..., x_n; \theta) = \prod_{i=1}^{n} f(x_i; \theta)$$
$$= f(x_1; \theta) \cdot f(x_2; \theta) \cdot ... \cdot f(x_n; \theta)$$

The log-likelihood function:

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

- Other notation: $\mathcal{L}(\theta|x_1, x_2, x_3, ..., x_n) = f(x_1, x_2, x_3, ..., x_n|\theta) = \prod_{i=1}^n f(x_i|\theta)$
- Example 1: A coin, with an unknown $\mathbb{P}(H) = p$, was flipped 10 times. The outcome is HHHTHHHTHH. What is the MLE for p?
- \circ 1. Specify the parameter $\theta:p$
- \circ 2. Specify the parameter space Ω : $\{p \mid 0 \leqslant p \leqslant 1\}$
- \circ 3. Write out the probability function $p_X(k)= \begin{cases} p & \text{, when } k=1\\ 1-p & \text{, when } k=0 \end{cases}$
- o 4. Write out the likelihood function:

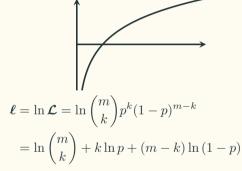
$$\mathcal{L}(p; 1110111011) = f(1110111011; p) = \prod_{i=1}^{10} f(x_i; p)$$

$$= f(1; p) \cdot f(1; p) \cdot f(1; p) \cdot f(0; p) \cdot f(1; p)$$

$$= p \cdot p \cdot p \cdot (1 - p) \cdot p \cdot p \cdot (1 - p) \cdot p \cdot p \cdot p \cdot (1 - p)^2$$

- Example 2 A more generalised case of coin flipping: A (possibly unfair) coin is flipped m times, and k heads are observed. Let $\mathbb{P}(H) = p$. What is the MLE for p?
- 1. $\theta : (n, p)$
- 2. Ω : { $(n,p) \mid n = m, 0 }$
- 3. $p_X(k) = \binom{n}{k} p^k (1-p)^{n-k}$
- 4.

$$\mathcal{L}(n,p;k) = f(k;n,p) = {m \choose k} p^k (1-p)^{m-k}$$



What value should p take to maximise ℓ ?

Let
$$\frac{\mathrm{d}\ell}{\mathrm{d}p} = 0 \ \Rightarrow \ \hat{p} = \frac{k}{m}$$

- Example 3 DNA synthesis errors: The genetic material is copied and synthesised by DNA polymerase. One high-fidelity DNA polymerase, Pfu, originally isolated from the hyperthermophilic archae Pyrococcus furiosus, is believed to have very low error rate. Assume the errors generated by Pfu follow a Poisson distribution with λ mutations per 10^6 base pairs (Mb). We have examined n newly synthesised DNA fragments and observed that the nubmer of mutations per Mb is $k_1, k_2, k_3, ..., k_n$. What is the MLE for λ ?
- 1. $\theta:\lambda$
- 2. $\Omega : \{ \lambda \mid \lambda > 0 \}$
- 3. $p_x(k) = \frac{\lambda^k}{k!}e^{-\lambda}$

- 4.
$$\mathcal{L}(\lambda; k_1, k_2, ..., k_n) = f(k_1, k_2, ..., k_n; \lambda) = \prod_{i=1}^n \frac{\lambda^{k_i}}{k_i!} e^{-\lambda}$$

- Example 4 Pou5f1 expression in embryonic stem cells: Let the random variable \boldsymbol{X} be the expression values of Pou5f1. We know $\boldsymbol{X} \sim \mathcal{N}(\mu, \sigma^2)$, but μ and σ^2 are unknown. Now we have sequenced n cells, and the expressions of Pou5f1 in those cells are $x_1, x_2, ..., x_n$, respectively. What is the MLE for the parameters of this normal distribution?
- 1. $\theta : \mu, \sigma^2$
- 2. $\Omega:\{(\mu,\sigma^2) \mid -\infty < \mu < +\infty, \ \sigma^2 \geqslant 0\}$
- 3. $f_X(x) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$
- 4. $\mathcal{L}(\mu, \sigma^2; x_1, x_2, ..., x_n) = f(x_1, x_2, ..., x_n; \mu, \sigma^2) = \prod_{i=1}^n \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x_i \mu)^2}{2\sigma^2}}$

Probability vs. Likelihood





the likelihood of the parameter(s) θ taking certain values given that a bunch of data $x_1, x_2, ..., x_n$ are observed.



the joint probability mass/density of observing the data $x_1, x_2, ..., x_n$ with model parameter(s) θ .

from Wolfram:

Likelihood is the hypothetical probability that an event that has already occurred would yield a specific outcome. The concept differs from that of a probability in that a **probability** refers to the occurrence of future events, while a **likelihood** refers to past events with known outcomes.

Advantages and Disadvantages of MLE

Advantages:

- Intuitive and straightforward to understand.
- If the model is correctly assumed, the MLE is efficient (meaning small variance or mean squared error).
- Can be extended to do other useful things.

Disadvantages:

- Relies on assumptions of a model (need to know the PMF/PDF).
- Sometimes difficult or impossible to solve the derivate of $\mathcal L$ or ℓ .
- Sometimes leads to the wrong or biased conclusions