Probability and Statistics 7 for Computer Science

"Statistical thinking will one day be as necessary for efficient citizenship as the ability to read and write." H. G. Wells

Credit: wikipedia

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Objectives

✺Hypothesis test

KEMaximum Likelihood Estimation

A hypothesis

✺ Ms. Smith's vote percentage is 55% This is what we want to test, often called null hypothesis H_0

✺ Should we reject this hypothesis given the poll data?

Fraction of "less extreme" samples

- Assuming the hypothesis H_0 is true
- Define a test statistic

$$
x = \frac{(sample\ mean) - (hypothesized\ value)}{standard\ error}
$$

 $*$ Since N>30, we assume x comes from a standard normal

 $*$ So, the fraction of "less extreme" samples is: $f =$ 1 $\sqrt{2\pi}$ $\int |x|$ $\int_{-|x|}^{|x|} exp(-\frac{u^2}{2}) du$

Rejection region of null hypothesis H_{o}

- Assuming the hypothesis H_0 is true
- Define a test statistic

 $x =$ $(sample \ mean) - (hypothesized \ value)$ standard error

Since N>30, assume x comes from a standard normal

P-value: Rejection region- "The extreme fraction"

 $*$ It is conventional to report the p-value of a hypothesis test

$$
p = 1 - f = 1 - \frac{1}{\sqrt{2\pi}} \int_{-|x|}^{|x|} exp(-\frac{u^2}{2}) du
$$

p-value: election polling

- $H₀$. Ms. Smith's vote percentage is 55%
- $*$ The sample mean is 51% and stderr is 1.44%
- $\frac{1}{2}$ The test statistic $x =$ $51 - 55$ 1.44 $=-2.7778$
- $*$ And the p-value for the test is:

$$
p = 1 - \frac{1}{\sqrt{2\pi}} \int_{-2.7778}^{2.7778} exp(-\frac{u^2}{2}) du = 0.00547 \qquad 0.05
$$

So we reject the hypothesis

Hypothesis test if $N < 30$

- $*$ Q: what distribution should we use to test the hypothesis of sample mean if N<30?
	- A. Normal distribution
	- B. t-distribution with degree $=30$
	- C. t-distribution with degree $= N$
	- D. t-distribution with degree $= N-1$

The use and misuse of p-value

- p-value use in scientific practice
	- ✺ Usually used to reject the null hypothesis that the data is random noise
	- $*$ Common practice is $p < 0.05$ is considered significant evidence for something interesting
- Caution about p-value hacking
	- \\ Rejecting the null hypothesis doesn't mean the alternative is true
	- $*$ P < 0.05 is arbitrary and often is not enough for controlling false positive phenomenon

Be wary of one tailed p-values

✺ The one tailed p-value should only be considered when the realized sample mean or differences will for sure fall only to one size of the distribution.

Sometimes scientist are tempted to use one tailed test because it'll give smaller p-val. But this is bad statistics!

Maximum likelihood estimation

The parameter estimation problem

- Suppose we have a dataset that we know comes from a distribution (ie. Binomial, Geometric, or Poisson, etc.)
- What is the best estimate of the parameters (**θ** or **θ**s) of the distribution?
	- Examples:
		- $*$ For binomial and geometric distribution, $\theta = p$ (probability of success)
		- $*$ For Poisson and exponential distributions, $\theta = \lambda$ (intensity)
		- $*$ For normal distributions, **θ** could be *μ* or $σ²$.

Motivation: Poisson example

 $*$ Suppose we have data on the number of babies born each hour in a large hospital

- $*$ We can assume the data comes from a Poisson distribution
- \mathscr{H} What is your best estimate of the intensity λ ?

Credit: David Varodayan

Maximum likelihood estimation (MLE)

 \mathscr{W} We write the probability of seeing the data D given parameter θ

$$
L(\theta) = P(D|\theta)
$$

 $\mathscr W$ The **likelihood function** $L(\theta)$ is **not** a probability distribution

θ is

EXECTE: The maximum likelihood estimate (MLE) of

$$
\hat{\theta} = arg \ max_{\theta} L(\theta)
$$

Why is $L(\theta)$ not a probability distribution?

A. It doesn't give the probability of all the $posible$ θ values.

B. Don't know whether the sum or integral of $L(\theta)$ for all possible θ values is one or not.

C. Both.

Likelihood function: Binomial example

- ✺ Suppose we have a coin with unknown probability of coming up heads
- **Example 18 We toss it N times and observe k heads**
- $*$ We know that this data comes from a binomial distribution
- **What** is the likelihood function $L(\theta) = P(D|\theta)$?

Likelihood function: binomial example

- ✺ Suppose we have a coin with unknown probability of coming up heads
- **Example 18 We toss it N times and observe k heads**
- $*$ We know that this data comes from a binomial distribution
- **What** is the likelihood function $L(\theta) = P(D|\theta)$?

$$
L(\theta) = {N \choose k} \theta^k (1-\theta)^{N-k}
$$

$$
L(\theta) = {N \choose k} \theta^k (1-\theta)^{N-k}
$$

 $\hat{\theta}$ In order to find: $\theta = arg\ max\ \overline{L}(\theta)$ We set: $dL(\theta)$ $\frac{\partial}{\partial \theta} = 0$

$$
L(\theta) = {N \choose k} \theta^k (1-\theta)^{N-k}
$$

$$
L(\theta) = {N \choose k} \theta^k (1-\theta)^{N-k}
$$

$$
\frac{d}{d\theta}L(\theta) = {N \choose k} (k\theta^{k-1}(1-\theta)^{N-k} - \theta^k(N-k)(1-\theta)^{N-k-1}) = 0
$$

$$
L(\theta) = {N \choose k} \theta^k (1-\theta)^{N-k}
$$

$$
\frac{d}{d\theta}L(\theta) = {N \choose k} (k\theta^{k-1}(1-\theta)^{N-k} - \theta^k(N-k)(1-\theta)^{N-k-1}) = 0
$$

$$
k\theta^{k-1}(1-\theta)^{N-k} = \theta^k(N-k)(1-\theta)^{N-k-1}
$$

$$
L(\theta) = \binom{N}{k} \theta^k (1-\theta)^{N-k}
$$

$$
\frac{d}{d\theta}L(\theta) = {N \choose k}(k\theta^{k-1}(1-\theta)^{N-k} - \theta^k(N-k)(1-\theta)^{N-k-1}) = 0
$$

$$
k\theta^{k-1}(1-\theta)^{N-k} = \theta^k(N-k)(1-\theta)^{N-k-1}
$$

$$
k - k\theta = N\theta - k\theta
$$

$$
L(\theta) = \binom{N}{k} \theta^k (1-\theta)^{N-k}
$$

$$
\frac{d}{d\theta}L(\theta) = {N \choose k} (k\theta^{k-1}(1-\theta)^{N-k} - \theta^k(N-k)(1-\theta)^{N-k-1}) = 0
$$

$$
k\theta^{k-1}(1-\theta)^{N-k} = \theta^k(N-k)(1-\theta)^{N-k-1}
$$

$$
k - k\theta = N\theta - k\theta
$$

$$
\widehat{\theta} = \frac{k}{N}
$$

The MLE of p

Likelihood function: geometric example

- ✺ Suppose we have a die with unknown probability of coming up six
- $\mathcal W$ We roll it and it comes up six for the first time on the kth roll
- \mathscr{W} We know that this data comes from a geometric distribution
- **What** is the likelihood function $L(\theta) = P(D|\theta)$? Assume $θ$ is p .

$$
L(\theta)=(1-\theta)^{k-1}\theta
$$

$$
L(\theta)=(1-\theta)^{k-1}\theta
$$

$$
\frac{d}{d\theta}L(\theta) = (1 - \theta)^{k-1} - (k - 1)(1 - \theta)^{k-2}\theta = 0
$$

$$
L(\theta)=(1-\theta)^{k-1}\theta
$$

$$
\frac{d}{d\theta}L(\theta) = (1 - \theta)^{k-1} - (k - 1)(1 - \theta)^{k-2}\theta = 0
$$

$$
(1 - \theta)^{k-1} = (k - 1)(1 - \theta)^{k-2}\theta
$$

$$
L(\theta)=(1-\theta)^{k-1}\theta
$$

$$
\frac{d}{d\theta}L(\theta) = (1 - \theta)^{k-1} - (k - 1)(1 - \theta)^{k-2}\theta = 0
$$

$$
(1 - \theta)^{k-1} = (k - 1)(1 - \theta)^{k-2}\theta
$$

$$
1 - \theta = k\theta - \theta
$$

$$
L(\theta)=(1-\theta)^{k-1}\theta
$$

$$
\frac{d}{d\theta}L(\theta) = (1 - \theta)^{k-1} - (k - 1)(1 - \theta)^{k-2}\theta = 0
$$

$$
(1 - \theta)^{k-1} = (k - 1)(1 - \theta)^{k-2}\theta
$$

$$
1 - \theta = k\theta - \theta
$$

$$
\hat{\theta}=\frac{1}{k}
$$

The MLE of p

MLE with data from IID trials

WE A If the dataset $D = \{x\}$ comes from IID trials

$$
L(\theta) = P(D|\theta) = \prod_{x_i \in D} P(x_i|\theta)
$$

 \mathcal{L} Each x_i is one observed result from an IID trial

Q: MLE with data from IID trials

WE A If the dataset $D = \{x\}$ comes from IID trials

$$
L(\theta) = P(D|\theta) = \prod_{x_i \in D} P(x_i|\theta)
$$

Why is the above function defined by the product?

A. IID samples are independent

B. Each trial has identical probability function

C. Both.

MLE with data from IID trials

WE A If the dataset $D = \{x\}$ comes from IID trials

$$
L(\theta) = P(D|\theta) = \prod_{x_i \in D} P(x_i|\theta)
$$

 $*$ The likelihood function is hard to differentiate in general, except for the binomial and geometric cases.

✺ Clever trick: take the (natural) log

Log-likelihood function

 $*$ Since log is a strictly increasing function

$$
\hat{\theta} = arg \; max \; L(\theta) = arg \; max \; logL(\theta)
$$

✺ So we can aim to maximize the **log-likelihood** function

$$
logL(\theta) = logP(D|\theta) = log \prod_{x_i \in D} P(x_i|\theta) = \sum_{x_i \in D} log P(x_i|\theta)
$$

 $*$ The log-likelihood function is usually much easier to differentiate

Log-likelihood function: Poisson example

 $*$ Suppose we have data on the number of babies born each hour in a large hospital

- $*$ We can assume the data comes from a Poisson distribution λ
- $\frac{1}{100}$ What is the log likelihood function $LogL(\theta)$?

Log-likelihood function: Poisson example

$$
L(\theta) = \prod_{i=1}^{N} \frac{e^{-\theta} \theta^{k_i}}{k_i!}
$$

MLE : Poisson example

$$
LogL(\theta) = \sum_{i=1}^{N} (-\theta + k_i log\theta - log k_i!)
$$
MLE : Poisson example

$$
LogL(\theta) = \sum_{i=1}^{N} (-\theta + k_i \log \theta - \log k_i!)
$$

$$
\frac{d}{d\theta} log L(\theta) = 0 \Rightarrow \sum_{i=1}^{N} (-1 + \frac{k_i}{\theta} - 0) = 0
$$

MLE : Poisson example

$$
LogL(\theta) = \sum_{i=1}^{N} (-\theta + k_i \log \theta - \log k_i!)
$$

$$
\frac{d}{d\theta} log L(\theta) = 0 \Rightarrow \sum_{i=1}^{N} (-1 + \frac{k_i}{\theta} - 0) = 0
$$

$$
-N + \frac{\sum_{i}^{N} k_i}{\theta} = 0
$$

MLE : Poisson example

$$
LogL(\theta) = \sum_{i=1}^{N} (-\theta + k_i \log \theta - \log k_i!)
$$

$$
\frac{d}{d\theta} log L(\theta) = 0 \Rightarrow \sum_{i=1}^{N} (-1 + \frac{k_i}{\theta} - 0) = 0
$$

$$
-N + \frac{\sum_{i}^{N} k_i}{\theta} = 0
$$

$$
\hat{\theta} = \frac{\sum_{i}^{N} k_{i}}{N}
$$

The MLE of λ

- **WE Suppose we model the dataset** $D = \{x\}$ as normally distributed
- $*$ What should be the likelihood function? Is the method of modeling the same as for the Poisson distribution?

A. Yes B. No

- **WE Suppose we model the dataset** $D = \{x\}$ as normally distributed
- $*$ What should be the likelihood function? Is the method of modeling the same as for the Poisson distribution? Yes and No. The idea is similar but the normal distribution is continuous, we need to use the **probability density** instead.

- **WE Suppose we model the dataset** $D = \{x\}$ as normally distributed
- $*$ The likelihood function of a normal distribution:

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

- **WE Suppose we model the dataset** $D = \{x\}$ as normally distributed
- ✺ There are two parameters to esGmate: *μ* and *σ* $*$ If we fix *σ* and set θ= *μ* $\hat{\bm{\rho}}$ $\hat{\theta} =$ 1 $\overline{\overline{N}}$ \sum \overline{N} $i=1$ \overline{x}_i
	- ✺ If we fix *μ* and set θ= *σ*

$$
\hat{\theta} = \sqrt{\frac{1}{N} \sum_{i=1}^{N} (x_i - \mu)^2}
$$

Drawbacks of MLE

- ✺ Maximizing some likelihood or log-likelihood function is mathematically hard
- ✺ If there are very few data items, the MLE estimate maybe very unreliable
	- **WE If** we observe 3 heads in 10 coin tosses, should we accept that $p(heads)=0.3$?
	- **WE If** we observe 0 heads in 2 coin tosses, should we accept that $p(heads)=0$?

Confidence intervals for MLE estimates

- $\mathscr K$ An MLE parameter estimate θ depends on the data that was observed $\overline{\hat{\hat{\mu}}}$ $\hat{\theta}$
- **We can construct a confidence interval for** $\hat{\theta}$ **using** the parametric bootstrap $\hat{\bm{\rho}}$ $\hat{\theta}$
	- **We Use the distribution with parameter** $\hat{\theta}$ to generate a large number of bootstrap samples $\hat{\theta}$
	- $*$ From each "synthetic" dataset, re-estimate the parameter using MLE
	- ✺ Use the histogram of these re-esGmates to construct a confidence interval

Assignments

✺ Finish Chapter 7 of the textbook

 $*$ Next time: Maximum likelihood estimate, Bayesian inference

Additional References

- ✺ Robert V. Hogg, Elliot A. Tanis and Dale L. Zimmerman. "Probability and Statistical Inference"
- ✺ Morris H. Degroot and Mark J. Schervish "Probability and Statistics"

Chi-square distribution

- $*$ If $Z_i's$ are independent variables of standard normal distribution, $\mathcal{L} = \mathcal{L}$ has a Chi-square distribution with degree of freedom \bm{m} , $X \sim \chi^2(m)$ $X = Z_1^2 + Z_2^2 + ... + Z_m^2 = \sum$ \overline{m} $i=1$ Z_i^2
- \mathscr{H} We can test the goodness of fit for a model using a statistic *C* against this distribution, where

$$
C = \sum_{i=1}^{m} \frac{(f_o(\varepsilon_i) - f_t(\varepsilon_i))^2}{f_t(\varepsilon_i)}
$$

Independence analysis using Chi-square

✺ Given the two way table, test whether the column and row are independent

Independence analysis using Chi-square

 $*$ The theoretical expected values if independent

The degree of the chi-square distribution for the two way table

✺ The degree of freedom for the chi-square distribution for a r by c table is

 $(r-1) \times (c-1)$ where $r>1$ and $c>1$

 $\frac{1}{2}$ Because the degree df = n-1-p See textbook Pg 171-172

 $=$ rc $-1-$ (r-1) $-$ (c-1)

n is the number of cells of $= (r-1) \times (c-1)$ p is the number of $= 2$ data;

unknown parameters

$$
= (r-1) \times (c-1)
$$

$$
=2
$$

Chi-square test for the popular kid data

 $*$ The Chi-statistic : 21.455

```
chisq.test(data_BG)
```
Pearson's Chi-squared test

```
data: data BG
X-squared = 21.455, df = 2, p-value = 2.193e-05
```
\n
$$
\text{Example: } \left| 2.193e-05 \right|
$$
\n

\nume\$\tequate line two categories are \\,\tequate line two categories are independent

Q. What is the degree of freedom for this?

$*$ The following 2-way table for chi-square test has a degree of freedom equal to:

Table 10.26 Data for Exercise 3

Chi-square test is very versatile

- $\mathcal K$ Chi-square test is so versatile that it can be utilized in many ways either for discrete data or continuous data via intervals
- ✺ Please check out the worked-out examples in the textbook and read more about its applications.

We are interested in comparing sample means

- Are the average daily body temperature of the two beavers the same?
- We need to model the difference between two sample means

How do we model the difference between two samples means?

- **E** We know when the sample size **N** is large, the sample mean random variable approaches normal $*$.
- **Example So our problem became finding the model** of the difference between two normally distributed random variables.

* Assume the daily temperature at different times are independent.

Background: sum of independent normals

$*$ We know

 $X_1 \sim normal(\mu_1, \sigma_1^2)$

 $X_2 \sim normal(\mu_2, \sigma_2^2)$

 $X_1 + X_2 \sim ?$ $\overline{\mathbf{?}}$

 \mathcal{K} The sum of X_1 and X_2 is still normal (proof omitted, ref. ...)

Background: sum of independent normals

✺ We know

 $X_1 \sim normal(\mu_1, \sigma_1^2)$

 $X_2 \sim normal(\mu_2, \sigma_2^2)$

$$
\text{# So} \quad X_1 + X_2 \sim \text{normal}(\mu_1 + \mu_2, \sigma_1^2 + \sigma_2^2)
$$

 $*$ By the linearity of expected value and the sum rule of variance of the sum of two independent random variables.

Background: sum of independent normals

 $*$ We know $X_1 \sim normal(\mu_1, \sigma_1^2)$

 $X_2 \sim normal(\mu_2, \sigma_2^2)$

 $\text{$ } **So** $\left| X_1 + X_2 \sim normal(\mu_1 + \mu_2, \sigma_1^2 + \sigma_2^2) \right|$

 $*$ By properties: $E[X_1+X_2]=E[X_1]+E[X_2]$ $var[X_1 + X_2] = var[X_1] + var[X_2]$

Difference of independent normals

✺ We know

$$
X_1 \sim normal(\mu_1, \sigma_1^2)
$$

$$
X_2 \sim normal(\mu_2, \sigma_2^2)
$$

$$
X_1 - X_2 \sim ?
$$

 $\mathscr K$ The difference of X_1 and X_2 is still normal (proof omitted)

Difference of independent normals

✺ We know $X_1 \sim normal(\mu_1, \sigma_1^2)$

$$
X_2 \sim normal(\mu_2, \sigma_2^2)
$$

$$
\ast \text{So} \quad \left[X_1 - X_2 \sim normal(\mu_1 - \mu_2, \sigma_1^2 + \sigma_2^2) \right]
$$

 $*$ By the linearity of expected value and the sum rule of variance of the sum of two independent random variables and the scaling property of variance.

✺ Because

✺ Because E[X¹ − X2] = E[X1] − E[X2] $= \mu_1 - \mu_2$

> $var[X_1 - X_2] = var[X_1 + (-X_2)]$ = variation of variation of variation.
The variation of va
Constitution of variation of variation of variation of variation of var

$$
var[X_1 - X_2] = var[X_1 + (-X_2)]
$$

= $var[X_1] + var[-X_2]$

$$
var[X_1 - X_2] = var[X_1 + (-X_2)]
$$

= $var[X_1] + var[-X_2]$
= $var[X_1] + var[X_2]$

$$
var[c \cdot X_2] = c^2 var[X_2]
$$

$$
var[X_1 - X_2] = var[X_1 + (-X_2)]
$$

= $var[X_1] + var[-X_2]$
= $var[X_1] + var[X_2]$
= $o_1^2 + o_2^2$

$$
var[X_1 - X_2] = var[X_1 + (-X_2)]
$$

= $var[X_1] + var[-X_2]$
= $var[X_1] + var[X_2]$
= $o_1^2 + o_2^2$

$$
\frac{\ast}{\left(X_1 - X_2 \sim normal(\mu_1 - \mu_2, \sigma_1^2 + \sigma_2^2)\right)}
$$

Now we are ready to check the differences between sample means

✺ Because sample means are roughly normal when *N* is large.

$$
\frac{\lambda_1 - \lambda_2 \sim normal(\mu_1 - \mu_2, \sigma_1^2 + \sigma_2^2)}{}
$$

The difference between two sample means

- $*$ Suppose we draw samples from two populations $\{x\}$ and $\{y\}$
	- **WE From a sample of size** k_x **from** $\{x\}$ **, we get sample** mean $X^{(k_x)}$
	- **WE From a sample of size** k_y **from** $\{y\}$ **, we get sample** mean $Y^{(k_y)}$

The difference between two sample means

- $*$ Define random variable $D = X^{(k_x)} Y^{(k_y)}$ as the difference between the sample means
- $*$ If we hypothesize that popmean($\{x\}$) = popmean($\{y\}$), then

$$
E[D] = E[X^{(k_x)}] - E[Y^{(k_y)}] = 0
$$

Standard error of the difference between two sample means

- $*$ Recall the standard error is roughly the standard deviation of a sample mean
- $\mathscr K$ By the property of variance of the difference between two independent normals

$$
var[D] \doteq stderr(\{x\})^2 + stderr(\{y\})^2
$$

\n
$$
std[D] \doteq \sqrt{stderr(\{x\})^2 + stderr(\{y\})^2} = stderr[D]
$$

\n
$$
std[D] \doteq \sqrt{\frac{stdunbiased(\{x\})^2}{k_x} + \frac{stdunbiased(\{y\})^2}{k_y}}
$$
P-value for testing the equality of two means

 $*$ Define the test statistic

$$
g = \frac{mean(\{x\}) - mean(\{y\})}{stderr(D)}
$$

 $*$ If k_x ≥ 30 and If k_y ≥ 30

$$
p = 1 - f = 1 - \frac{1}{\sqrt{2\pi}} \int_{-|g|}^{|g|} exp(-\frac{u^2}{2}) du
$$

P-value: Rejection region- "The extreme fraction"

 $*$ It is conventional to report the p-value of a hypothesis test

$$
p = 1 - f = 1 - \frac{1}{\sqrt{2\pi}} \int_{-|g|}^{|g|} exp(-\frac{u^2}{2}) du
$$

Since *N*>30, *x* should come from a standard normal

Comparing the body temperatures of two beavers

$$
k_{x} = 114 \text{ and } k_{y} = 100
$$
\n
$$
k_{x} = 114 \text{ and } k_{y} = 100
$$
\n
$$
k_{y} = 100
$$
\n
$$
k_{x} = 114 \text{ and } k_{y} = 100
$$
\n
$$
k_{y} = 100
$$
\n<

Comparing the body temperatures of two beavers

WE Hypothesis H₀: the mean temperatures of the two beavers are the same

Example 11 The test statistic
$$
g = \frac{36.86219 - 37.5967}{0.04821181} = -15.235
$$

$$
p = 1 - f = 1 - \frac{1}{\sqrt{2\pi}} \int_{-15.235}^{15.235} exp(-\frac{u^2}{2}) du
$$

$$
p \simeq 0
$$

 $*$ So we can reject the hypothesis that the mean temperatures are the same

What if $N < 30$?

- $*$ There are general solutions for either N $>=$ 30 or N $<$ 30 if the data sets are random samples from normal distributed data.
	- ✺ The difference between sample means can be either modeled as t-distribution with degree (k_r) $+k_y-2$) when their population standard deviations are the same
	- ✺ Or the difference between sample means can be approximated with t-distribution with other proper degree of freedom.
	- ✺ There are build in t-test procedures in Python, R

Compare the two mean temperatures of two beavers with t.test

WE Hypothesis H₀: the mean temperatures of the two beavers are the same

> t.test(beaver1\$temp, beaver2\$temp)

Welch Two Sample t-test

```
data: beaver1$temp and beaver2$temp
t = -15.235, df = 131.12, p-value < 2.2e-16
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -0.8298806 - 0.6391334sample estimates:
mean of x mean of y
 36.86219 37.59670
```
$*$ $p < 2.2e-16$, also reject the hypothesis

See you next time

See You!

