TIARA Summer School on Astrostatistics: Statistical Inference

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- Statistical inference is about how we use data to infer the underlying population that generates our data.
- This process often involves a statistical model.
- Given a data (sample), a statistical model is a probability distribution that describes how this data is generated.
- Using the concept of statistical models, we can simply say that statistical inference is how we use the *observed data* to infer some characteristics of the *unobserved (probability) distribution*.

- What will the data look like?
- Here is part of the data in the SDSS: It is about 100 galaxy's log stellar mass:

 $\begin{array}{c} 11.26 \ 10.76 \ 11.57 \ 11.12 \ 11.25 \ 11.29 \ 11.32 \ 11.46 \ 10.93 \ 11.34 \ 9.12 \ 11.09 \ 11.11 \\ 11.62 \ 11.06 \ 11.22 \ 10.94 \ 11.33 \ 10.45 \ 11.79 \ 11.01 \ 11.40 \ 11.38 \ 11.16 \ 11.19 \ 11.47 \\ 11.38 \ 11.24 \ 11.05 \ 11.43 \ 11.26 \ 11.12 \ 11.24 \ 11.20 \ 11.55 \ 11.43 \ 11.22 \ 11.36 \ 11.38 \\ 11.27 \ 11.04 \ 11.72 \ 11.27 \ 11.16 \ 10.85 \ 11.45 \ 11.37 \ 11.17 \ 11.25 \ 11.10 \ 11.27 \ 11.41 \\ 11.15 \ 11.43 \ 11.22 \ 11.61 \ 11.34 \ 11.64 \ 11.53 \ 11.26 \ 11.19 \ 11.20 \ 11.20 \ 11.20 \ 11.52 \ 10.49 \\ 11.18 \ 11.19 \ 11.52 \ 11.32 \ 11.46 \ 11.03 \ 11.43 \ 11.26 \ 11.13 \ 11.32 \ 11.92 \ 10.94 \ 11.29 \\ 11.58 \ 11.11 \ 11.25 \ 11.69 \ 11.28 \ 11.40 \ 11.33 \ 11.26 \ 11.31 \ 11.32 \ 11.92 \ 10.94 \ 11.29 \\ 11.60 \ 11.26 \ 11.15 \ 11.43 \ 11.26 \ 11.28 \ 11.41 \\ 11.60 \ 11.26 \ 11.16 \ 11.28 \ 11.43 \ 11.26 \ 11.28 \ 11.46 \\ 11.30 \ 11.26 \ 11.26 \ 11.28 \ 11.18 \\ 11.60 \ 11.26 \ 11.17 \ 11.35 \ 11.43 \ 11.26 \ 11.22 \ 11.66 \\ 11.26 \ 11.26 \ 11.26 \ 11.38 \ 11.44 \ 11.31 \ 11.22 \ 11.28 \ 11.88 \\ 11.60 \ 11.26 \ 11.17 \ 11.35 \ 11.43 \ 11.26 \ 11.22 \ 11.66 \\ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.41 \\ 11.60 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.28 \ 11.48 \\ 11.60 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.28 \ 11.48 \\ 11.60 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.28 \ 11.48 \\ 11.60 \ 11.26 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.48 \\ 11.60 \ 11.26 \ 11.26 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.26 \ 11.28 \ 11.46 \ 11.26 \ 11.26 \ 11.28 \ 11.44 \ 11.26 \ 11.26 \ 11.28 \ 11.46 \ 11.44 \ 11.44 \ 11.46 \$

- Often our data is just a collection of numbers.
- A statistical model is a distribution that generates these numbers.

- In statistics, the values of our data are viewed as random variables X₁, · · · , X_n (n : sample size) that are IID from a distribution P(x).
- In most cases, we will further assume that such a distribution P(x) has a probability density function p(x).
- The above procedure will often be simply written as

$$X_1,\cdots,X_n\sim P$$

or

$$X_1,\cdots,X_n\sim p.$$

Parameters and Statistics

- Parameters (of interest): numbers or quantities that are features of the population distribution/density.
 - Examples: mean, median, mode, standard deviation (SD), regression coefficients, ···
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 - Example: sample average, sample median, sample SD, estimated regression coefficients, ···
 - From a mathematical point of view, a statistic is a function of random variables.
- Estimators: when a statistic is used to estimate a parameter, then this statistic is called an estimator (of the corresponding parameter).

Big Picture of Statistical Inference



Estimators and Estimation Theory

Estimating Basic Parameters

- Some parameters have a simple statistic that correspond to each of them.
- If we are interested in estimating these parameters, we can use these simple statistics.
- Example:

sample mean \longleftrightarrow population mean sample median \longleftrightarrow population median sample SD \longleftrightarrow population SD

Parametric Model - 1

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- Namely, the density function (or distribution function) is

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- Namely, the density function (or distribution function) is

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for some parameter θ .

In parametric model, the parameter θ completely determines the distribution so they are often the parameters of interest.

Parametric Model - 2

Here are some examples of parametric models:

• Normal distribution (parameters: μ and σ):

$$p(x;\mu,\sigma) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma}}$$

Exponential distribution (parameter: λ):

$$p(x;\lambda) = \lambda e^{-\lambda x}, x \ge 0.$$

Bernoulli distribution (parameter: p):

$$P(X = 1) = p$$
, $P(X = 0) = 1 - p$.

• Poisson distribution (parameter: λ):

$$P(X=k)=\frac{\lambda^k}{k!}e^{-\lambda}, k=0,1,2,\cdots.$$

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- For a given parameter θ, according to the parametric model p(x) = p(x; θ), the probability density generating X₁ is p(X_i; θ).
- Then we ask a simple question: for all possible values of the parameter θ, which value has the highest probability density of generating X₁?

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The MLE is then be written as

$$\widehat{ heta}_{MLE} = \mathop{argmax}\limits_{ heta} L(heta|X_i).$$

► In the case of observing *n* points, X₁, · · · , X_n, the likelihood function is

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Note that in the case where the random variables are discrete random variables (such as the Bernoulli model or Poisson model), we will replace the density function by the probability mass function.

▶ In the case of normal distribution, you can find that the MLE $\widehat{\mu}_{\textit{MLE}}$ and $\widehat{\sigma}_{\textit{MLE}}^2$ are

$$\widehat{\mu}_{MLE} = \overline{X}_n = \frac{1}{n} \sum_{i=1}^n, \quad \widehat{\sigma}_{MLE}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \overline{X}_n)^2 = \frac{n-1}{n} S_n^2.$$

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- The MLE of mean parameter is the sample mean and the MLE of SD parameter is similar to the sample SD.
- Note that the parameter p in a Bernoulli distribution and the parameter λ in a Poisson distribution both have the same form of MLE: the sample mean.
- The MLE of an exponential distribution is a bit more interesting.

- ▶ Recall that an exponential distribution has a probability density function $p(x; \lambda) = \lambda e^{-\lambda x}$.
- ▶ Thus, after observing X_1, \dots, X_n , the likelihood function is

$$L(\lambda|X_1,\cdots,X_n)=\prod_{i=1}^n\lambda e^{-\lambda X_i}=\lambda^n e^{-\lambda\sum_{i=1}^n X_i}.$$

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 Because taking the logarithm will not affect the position of maximum, we take the log of it. This leads to the log-likelihood function:

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$$\blacktriangleright \text{ Thus, } \widehat{\lambda}_{MLE} = \frac{n}{\sum_{i=1}^{n} X_i} = \frac{1}{\overline{X}_i}.$$

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- Let θ be the parameter of interest and $\hat{\theta}_n$ be an estimator for it.
- How do we quantify the accuracy of the estimator?
- ▶ Beware: since the estimator is computed from the data, the randomness of data will propagate to $\hat{\theta}_n$. So $\hat{\theta}_n$ is often a random quantity.

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- ► To quantify the accuracy of $\hat{\theta}_n$, we introduce two measures: bias and variance.
- The bias of an estimator is the systematic deviation from its target. Mathematically, we define it as

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- The bias of an estimator is the systematic deviation from its target. Mathematically, we define it as

$$\mathsf{bias}(\widehat{\theta}_n) = \mathbb{E}(\widehat{\theta}_n) - \theta.$$

► The variance describes the amount of randomness that an estimator has. It is simply the quantity Var(\(\heta_n\)), variance of the estimator.

- A: large bias, small variance.
- B: small bias, large variance.
- C: large bias, large variance.
- D: small bias, small variance.
- Ideally, we want an estimator with small bias and small variance (case D).



Estimation Theory: MSE

- An estimator is called *consistent* if when the sample size $n \to \infty$, $\hat{\theta}_n$ converges to θ in probability.
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Estimation Theory: MSE

- An estimator is called *consistent* if when the sample size $n \to \infty$, $\hat{\theta}_n$ converges to θ in probability.
- ▶ Both the bias and variance converges to 0 ⇒ it is a consistent estimator.
- There is a simple error measurement that takes into account both the bias and variance called the *mean square error (MSE)*.
- The MSE is

$$MSE(\widehat{\theta}_n, \theta) = \mathbb{E}\left((\widehat{\theta}_n - \theta)^2\right) = bias^2(\widehat{\theta}_n) + Var(\widehat{\theta}_n).$$

 The last equality is also known as the bias-variance decomposition.
Estimation Theory: Some Remarks

- A good news: most MLE's are consistent, so are the simple estimators of basic parameters.
- The consistency of an estimator depends on the assumptions about the population distribution.
- Even we are using the same estimator, it might be consistent for one dataset but inconsistent for another.
- An inconsistent estimator may lead you to a wrong conclusion.

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- A CI requires a number called confidence level, often denoted as 1 − α, and a CI is an *interval* C_{α,n} = [L_{α,n}, U_{α,n}] that can be computed using the data with the following property:

$$P(\theta \in C_{\alpha,n}) = P(L_{\alpha,n} \le \theta \le U_{\alpha,n}) \approx 1 - \alpha.$$

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- The randomness in the above probability comes from the randomness of L_{α,n}, U_{α,n} not θ!
- The lower bound L_{α,n} and upper bound U_{α,n} are statistics (numbers computed from the data).

Confidence Interval: an Illustration



Confidence Interval: Bernoulli Distribution

- ► The plot in the previous slide shows an example for a ≈ 95% CI for inferring the parameter p in a Bernoulli distribution.
- The interval with

$$L_n = \widehat{p} - 2\sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}}, U_n = \widehat{p} + 2\sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}}, \widehat{p} = \frac{1}{n}\sum_{i=1}^n X_i$$

is a $\approx 95\%$ confidence interval for the parameter *p*.

• Note that $\hat{p} = \frac{1}{n} \sum_{i=1}^{n} X_i$ is the MLE for the parameter p.

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- Note that $\hat{p} = \frac{1}{n} \sum_{i=1}^{n} X_i$ is the MLE for the parameter p.
- Common CIs have three components as indicated by the colors:
 - The estimator.
 - Standard error (SE) of the estimator.
 - Multiplier: determined by 1α , the confidence.

Standard Error of an Estimator

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- SE is the rough size of the error of our estimator.
- When reading papers, people often report the estimated value and its error – this error is the SE.
- For the estimator \hat{p} in the Bernoulli model, its variance

$$Var(\hat{p}) = Var\left(\frac{1}{n}\sum_{i=1}^{n}X_{i}\right)$$

$$= \frac{1}{n^{2}}\sum_{i=1}^{n}Var(X_{i}) \quad (independence)$$

$$= \frac{1}{n}Var(X_{1}) \quad (identical)$$

$$= \frac{p(1-p)}{n}.$$

$$\bullet \text{ Thus, the SD of } \hat{p} \text{ is } \sqrt{\frac{p(1-p)}{n}}, \text{ which can be approximated by}$$

$$\sqrt{\frac{\hat{p}(1-\hat{p})}{n}}.$$

- A multiplier is a number depending on the distribution of the estimator.
- Thanks to the central limit theorem, many estimators will be normally distributed around their targeted parameters.
- ► Namely, if we use $\hat{\theta}_n$ to estimate θ , under good assumptions we have

$$\widehat{\theta}_n \approx N(\theta, SE^2(\widehat{\theta}_n)).$$

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This implies

$$\frac{\widehat{\theta}_n - \theta}{SE(\widehat{\theta})} \approx N(0, 1).$$

- Let $z_{1-\alpha/2}$ be the number such that $P(|N(0,1)| \le z_{1-\alpha/2}) = 1 \alpha$.
- Then we have:

$$P\left(\left|\frac{\widehat{\theta}_n-\theta}{SE(\widehat{\theta})}\right|\leq z_{1-\alpha/2}\right)\approx 1-\alpha.$$

► This fact:

$$P\left(\left|\frac{\widehat{\theta}_n-\theta}{\mathsf{SE}(\widehat{\theta})}\right|\leq z_{1-\alpha/2}\right)\approx 1-\alpha.$$

implies that

$$P\left(\widehat{\theta}_n - z_{1-\alpha/2} \cdot SE(\widehat{\theta}_n) \le \theta \le \widehat{\theta}_n + z_{1-\alpha/2} \cdot SE(\widehat{\theta}_n)\right) \approx 1-\alpha.$$

Namely, $\widehat{\theta}_n \pm z_{1-\alpha/2} \cdot SE(\widehat{\theta}_n)$ is a $1-\alpha$ CI of θ .

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- ▶ Namely, $\hat{\theta}_n \pm z_{1-\alpha/2} \cdot SE(\hat{\theta}_n)$ is a 1α Cl of θ .
- For a normal distribution, $z_{0.975} \approx 1.96 \approx 2$ ($\alpha = 0.05 = 5\%$).

▶ By identifying
$$\hat{\theta}_n = \hat{p}$$
 and $SE(\hat{\theta}_n) = SE(\hat{p}) \approx \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$ and $z_{0.975} \approx 2$, we conclude that $\hat{p} \pm 2\sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$ is a CI of the parameter p in the Bernoulli distribution.

Confidence Interval: Mean

- ► Assume that we observe X₁, · · · , X_n ~ P and we are interested in the population mean µ.
- A simple estimator of the mean is the sample mean $\bar{X}_n = \frac{1}{n} \sum_{i=1}^n X_i$.
- The SD of the sample mean is

$$\sqrt{\operatorname{Var}(\bar{X}_n)} = \sqrt{\frac{\sigma^2}{n}},$$

where σ^2 is the variance of the population distribution (population variance).

- We can simply estimate the population variance by sample variance S²_n = ¹/_{n-1} ∑ⁿ_{i=1}(X_i − X̄_n)².
- Thus, a 95% CI of the mean μ is

$$\left[\bar{X}_n - 1.96 \cdot \frac{S_n}{\sqrt{n}}, \ \bar{X}_n + 1.96 \cdot \frac{S_n}{\sqrt{n}}\right].$$

Confidence Interval: Example

- Now going back to the galaxy example mentioned at the beginning. We have n = 100 galaxies.
- After computation, we found that the sample mean of log stellar mass $\bar{X}_n = 11.25$ and sample SD is 0.31.
- ▶ Thus, the SE of the sample mean is about $\frac{0.31}{\sqrt{100}} = 0.031$
- A 95% Cl of the mean log stellar mass is

 $[11.25 - 1.96 \cdot 0.031, 11.25 + 1.96 \cdot 0.031] = [11.19, 11.31].$

Confidence Interval: Some Remarks

- The significance level is often chosen by the researcher.
- ► Common choices are 95%, 90%, and 99% (α = 0.05, 0.1, 0.01).
- The corresponding number $z_{1-\alpha/2}$ will be

 $z_{0.975} \approx 1.96, \ z_{0.95} \approx 1.64, \ z_{0.996} \approx 2.58.$

- Note that a CI can also be one-sided. Namely, it can also be an interval like (−∞, c] or [c, ∞) for some constant c.
- Often the construction of CIs depends on the (asymptotic/limiting) distribution of the estimator. Normal distribution is a common case but sometimes people will use other distributions such as *t*-distribution, χ²-distribution, *F*-distribution, etc.

- Hypothesis test is a statistical procedure to make inference.
- Actually, many scientific discoveries implicitly or explicitly used the hypothesis test.
- In hypothesis test, we are comparing two hypothesis:
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 - Null hypothesis (H_0) : a statement we are testing its validity.
 - ► Alternative hypothesis (*H_a*): a statement against to the null hypothesis.
- In scientific research, the alternative hypothesis is often something we want to prove (using data) – we will explain this later.

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 - 1. We design a test statistic T_n (can be computed from the data).
 - 2. We study the distribution of such a test statistic assuming H_0 is true.
 - 3. Using the data, we then compute the value of the observed test statistics T_n .

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- Here is a brief description of the testing procedure.
 - 1. We design a test statistic T_n (can be computed from the data).
 - 2. We study the distribution of such a test statistic *assuming* H_0 *is true*.
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 - 5. We reject H_0 if P-value is smaller than the significance level α .

- Again we use the galaxy stellar mass data as an example.
- Assume that the previous literature suggested that the log stellar mass is

$$H_0: \mu = 11.15.$$

We want to test if this statement is reasonable using our data. Note that in this case, the alternative hypothesis is

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► Then we need to find the distribution of this test statistic. And it turns out that such a test statistic T_n, under H₀, has a nice distribution:

$$T_n \approx N(0,1).$$

Using the data, we compute the observed value of test statistic:

$$t_n = \frac{11.25 - 11.15}{0.031} = 3.23.$$

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If we choose a significance level of 5%, 1%, we will all reject H₀. In this case, we will claim that we have strong evidence to reject H₀ : µ = 11.25 under a significance level of 5% (or 1%).

Hypothesis Test: Significance Level

- ► The significance level can be interpreted as a tolerance level of wrongly reject H₀ (later we will call it type-1 error rate).
- ► If H₀ is correct, then the distribution of P-value will be a uniform distribution over 0 and 1 (you can try to prove this).

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- So when H₀ is correct, the chance of rejecting H₀ under a significance level α is α.
- A lower value of α requires a stronger evidence against H_0 to reject it.
- ▶ In our case, we can reject H_0 under $\alpha = 5\%, 1\%$ but not 0.1%.
- ► Rejecting H₀ implies that the claim µ = 11.15 is not reasonable so we conclude µ ≠ 11.15.
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- ▶ Proof by contradiction: if we want to prove a statement S, we assume that statement to be incorrect first and then show that this leads to a contradiction.
- ► In the case of hypothesis test, we are doing a very similar job: we first assume H₀ being correct and then show that H₀ contradicts to our data.
- The P-value is a quantity that serves as a measure of consistency between H₀ and our data. Thus, a low P-value means that H₀ is not consistent with data (i.e., they contradict to each other) so we reject the null hypothesis.
- Thus, you can easily see that the alternative hypothesis will be something we want to prove (because it is the complement of the null hypothesis).

Hypothesis Test: Type-1 and Type-2 Error

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- Type-1 error: the H_0 is correct but we mistakenly reject H_0 .
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- This implies: we are controlling type-1 error rate to be small in hypothesis test framework.
- Why do we want to control type-1 error rate? → We are doing a 'proof by contradiction' so we want to make sure we have strong evidence to 'prove' that H₀ is incorrect.
- ▶ Rejecting under a very small α. ⇔ Type-1 error rate is very small. ⇔ We have very strong evidence.

Hypothesis Test: Other Examples

Sometimes the null hypothesis will be a one-sided case such as

$$H_0: \mu \le 11.15.$$

In this case, the calculation of P-value will be slightly different since the 'more extreme' case will be on the other side.

▶ When we assume the parametric model is correct, many null hypothesis will be about the value of parameter. For instance, $X_1, \dots, X_m \sim N(0, \sigma^2)$ and

$$H_0: \sigma^2 = 0.2, \quad H_a: \sigma^2 \neq 0.2.$$

The hypothesis test and CI have a close relationship. In some cases, you can use one to compute the other.

Two-Sample Test - 1

- Two-sample test is a very important topic in scientific research.
- The goal of a two-sample test is to see if we have strong evidence that the two observed samples are from different populations.
- For instance, in analyzing galaxies, we can separate galaxies into two groups: elliptical galaxies and spiral galaxies. The two-sample test can be used to check if the distributions of stellar mass of the two populations are the same or not.

Two-Sample Test - 2

- Let X_1, \dots, X_n and Y_1, \dots, Y_m be the two samples we observed.
- Using statistical models, we model that

$$X_1, \cdots, X_n \sim P_X, \qquad Y_1, \cdots, Y_m \sim P_Y,$$

where P_X and P_Y are the distributions generating the two samples.

The two-sample test examines the following hypothesis:

$$H_0: P_X = P_Y$$

against

$$H_a: P_X \neq P_Y.$$

- A simple approach of the two-sample test is the mean test.
- Because

$$H_0: P_X = P_Y$$

implies $\mu_X = \mu_Y$ (μ_i is the mean of P_i), the mean test is to test

$$H_0: \mu_X = \mu_Y.$$

• Testing $\mu_X = \mu_Y$ is equivalent to testing

$$H_0: \mu_X - \mu_Y = 0.$$

So the test statistics is to use the difference between sample means X
n and Y
m and rescale it by the variance.

The sample means have variance

$$\operatorname{Var}(\bar{X}_n) = rac{\sigma_X^2}{n}, \quad \operatorname{Var}(\bar{Y}_m) = rac{\sigma_Y^2}{m},$$

where σ_X^2 and σ_Y^2 are the variance of P_X and P_Y .

• Thus, the quantity $\bar{X}_n - \bar{Y}_m$ has variance $\frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}$ (because they are independent).

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- Thus, the quantity $\bar{X}_n \bar{Y}_m$ has variance $\frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}$ (because they are independent).
- ▶ Because we do not know σ_X^2 and σ_Y^2 in practice, we will replace them by the sample variance S_X^2 and S_Y^2 .

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- ▶ Because we do not know σ_X^2 and σ_Y^2 in practice, we will replace them by the sample variance S_X^2 and S_Y^2 .
- Thus, our final test statistics is

$$T_{n,m} = \frac{\bar{X}_n - \bar{Y}_m}{\sqrt{\frac{S_X^2}{n} + \frac{S_Y^2}{m}}}$$

• $T_{n,m}$ will follow asymptotically a standard normal distribution (think about why) so we can compare $T_{n,m}$ to the standard normal to obtain a p-value.

- Sometimes people will compare T_{n,m} to a t-distribution rather than the standard normal distribution. This is because when both samples are from normal distributions (with different means), T_{n,m} has an exact distribution that is the t-distribution.
- ▶ When using *t*-distribution, such a test is called a T-test.
- When using a standard normal distribution, this test is called a Z-test.
- In addition to testing the mean, one can also test other parameters such as median and variance.

Nonparametric Method: KS-test – 1

- ► Here we introduce a famous test that directly test H₀ : P_X = P_Y - the KS-test.
- The KS-test (Kolmogorov-Smirnov test) is a classical approach in nonparametric two-sample test.

Nonparametric Method: KS-test – 1

- ► Here we introduce a famous test that directly test H₀ : P_X = P_Y - the KS-test.
- The KS-test (Kolmogorov-Smirnov test) is a classical approach in nonparametric two-sample test.
- ► We can estimate the distribution function P_X(x) by the empirical distribution function (EDF):

$$\widehat{P}_X(t) = \frac{1}{n} \sum_{i=1}^n I(X_i \le t),$$

where I(x) is the indicator function such that if the input is true, then it outputs 1 otherwise 0.

► Actually, P_X(t) is the ratio of data points X₁, · · · , X_n whose value is less than or equal to t.

Nonparametric Method: KS-test - 2

Here is an example of the EDF of 5 observations of 1, 1.2, 1.5, 2, 2.5:



Nonparametric Method: KS-test - 3

EDF (black curve) versus the true CDF of a standard normal distribution:



Nonparametric Method: KS-Test - 4

The KS-test is to use the following test statistics:

$$K_{n,m} = \sup_{t} |\widehat{P}_X(t) - \widehat{P}_Y(t)|,$$

where \sup_t is a mathematical generalization of \max_t (you can just view it as taking the maximum).

- ► After rescaling, the test statistics *K*_{*n*,*m*} has a known limiting distribution called the *Kolmogorov distribution*.
- ► An appealing feature is that the Kolmogorov distribution does not depend on the true distribution P_X and P_Y.
- We then reject the null hypothesis when $K_{n,m}$ is sufficiently large.

Two-Sample Test: Remarks

- The tests we mentioned previously are just common approaches of two-sample test.
- There are many other approaches if you are interested in, you can search permutation test, rank test, signed-rank test.
- You can even use histogram to do a two-sample test.
- Keep in mind: there is no universal optimal test and every test works under different assumptions.

- When we have a theoretical result and we want to compare our data to the theoretical result, we can use the goodness-of-fit test.
- A simple approach to achieve this is through the χ^2 test¹.

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- When we have a theoretical result and we want to compare our data to the theoretical result, we can use the goodness-of-fit test.
- A simple approach to achieve this is through the χ^2 test¹.
- For instance, after some computations, we may obtain the following table (numbers inside parentheses are the SE's):

Value (Errors)	Case 1	Case 2	Case 3	Case 4	Case 5
Observed	16.5 (0.5)	22.1 (0.3)	27.7 (2.2)	25.5 (0.5)	13.2 (0.4)
Theory	15	23	31	25	10

Can we make some conclusions about the theory using observed data?

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- In the above example, we have 5 statistics X₁, · · · , X₅ (observed value) and each of them has error σ₁, · · · , σ₅.
- We use μ_1, \cdots, μ_5 to denote the theoretical result.
- Goal: we want to see if our data fits to the theoretical calculations.
- The χ^2 -test compute the test statistic

$$T = \left(\frac{X_1 - \mu_1}{\sigma_1}\right)^2 + \dots + \left(\frac{X_5 - \mu_5}{\sigma_5}\right)^2$$

- ► If the theory is correct and the noises are independent and Gaussian, then the test statistic *T* follows a *χ*²₅, a *χ*²-distribution with 5 degrees of freedom.
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- After computing T = t from the data, the corresponding P-value is P(χ²_ν ≥ t) for ν degrees of freedom.

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- After computing T = t from the data, the corresponding P-value is P(χ²_ν ≥ t) for ν degrees of freedom.
- Note that the null hypothesis being test is

$$H_0: X_i \sim N(\mu_i, \sigma_i^2), i = 1, \cdots, \nu.$$

Recall the observed table:

Value (Errors)	Case 1	Case 2	Case 3	Case 4	Case 5
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 \blacktriangleright Thus, in this case, the χ^2 statistic is

$$T = \left(\frac{X_1 - \mu_1}{\sigma_1}\right)^2 + \dots + \left(\frac{X_5 - \mu_5}{\sigma_5}\right)^2$$

= $\left(\frac{16.5 - 15}{0.5}\right)^2 + \left(\frac{22.1 - 23}{0.3}\right)^2$
+ $\left(\frac{27.7 - 31}{2.2}\right)^2 + \left(\frac{25.5 - 25}{0.5}\right)^2 + \left(\frac{13.2 - 10}{0.4}\right)^2$
= $3^2 + 3^2 + 1.5^2 + 1^2 + 8^2$
= 85.25

 Thus, we should report that we observed a signal of a 85.25 χ² statistic with 5 degrees of freedom.

Goodness-of-fit Test: Remarks

- When the χ² statistic is large, it means that the data contradicts to the theory or the assumptions about the noises.
- ► When the \(\chi^2\) statistic is small, it means that the data seems to fit to the theory BUT this does NOT imply that the theory is correct!

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- Because the null hypothesis is the theory being true, the goodness-of-fit test can be used to 'prove' that the theoretical result is wrong but it CANNOT be used to prove that the theory is correct!
- Just think about the proof by contradiction: if we cannot show a statement contradicts to itself, this does not imply that statement to be true.

Correlation Test and Independence Test

 There are methods to test if two random variables are correlated. Namely, testing

 $H_0: \operatorname{corr}(X, Y) = 0,$

where corr(X, Y) is the correlation between random variable X and Y.

 Also, we can even test if two random variables are independent. Namely, testing

 $H_0: X \text{ and } Y \text{ are independent.} (\Leftrightarrow p(x, y) = p(x)p(y)).$

One can use Energy statistics² or the approach from reproducing kernel Hilbert space³.

²https://en.wikipedia.org/wiki/Energy_distance#Energy_statistics

³http://www.kyb.mpg.de/fileadmin/user_upload/files/publications/attachments/ NIPS2007-Gretton_%5b0%5d.pdf

Assume that we are doing hypothesis test 100 times with type-1 errors all being 5%, then it is very likely we are going to reject some H₀'s even when they are correct!

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- ► To avoid falsely rejecting some H₀, we often try to control the Familywise Error Rate (FWER): the chance of falsely rejecting any H₀.
- ► To make sure FWER is less than α, a classical approach is to use the Bonferroni correction⁴: we only reject those H₀ if their individual p-value is less than α/K where K is the total number of null hypothesis being tested.

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- There is another criterion that many people commonly use: instead of controlling the FWER, we control the False Discovery Rate (FDR)⁵.

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5 https://en.wikipedia.org/wiki/False discovery rate

Useful References

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