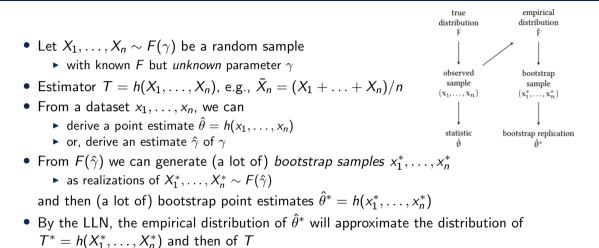
### Statistical Methods for Data Science Lesson 20 - Parametric bootstrap. Hypotheses testing.

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### Parametric bootstrap principle



### Parametric bootstrap

PARAMETRIC BOOTSTRAP SIMULATION (FOR  $\bar{X}_n - \mu$ ). Given a dataset  $x_1, x_2, \ldots, x_n$ , compute an estimate  $\hat{\theta}$  for  $\theta$ . Determine  $F_{\hat{\theta}}$  as an estimate for  $F_{\theta}$ , and compute the expectation  $\mu^* = \mu_{\hat{\theta}}$  corresponding to  $F_{\hat{\theta}}$ .

- 1. Generate a bootstrap dataset  $x_1^*, x_2^*, \ldots, x_n^*$  from  $F_{\hat{\theta}}$ .
- 2. Compute the centered sample mean for the bootstrap dataset:

$$\bar{x}_n^* - \mu_{\hat{\theta}}$$

where

$$\bar{x}_n^* = \frac{x_1^* + x_2^* + \dots + x_n^*}{n}$$

Repeat steps 1 and 2 many times.

- Use the empirical distribution of  $\delta^* = \bar{x}^*_n \mu_{\hat{\theta}}$  for estimating
  - ► confidence interval  $(c_l, c_u)$  for  $\delta = \bar{x}_n \mu$  as  $(q_{\alpha/2}, q_{1-\alpha/2})$  of  $\delta^*$  distribution

• 
$$c_l \leq \delta = \bar{x}_n - \mu \leq c_u$$
 implies  $\bar{x}_n - c_u \leq \mu \leq \bar{x}_n - c_l$ , i.e. c.i. for  $\mu$  is  $(\bar{x}_n - c_u, \bar{x}_n - c_l)$ 

#### See R script

## Application: distribution fitting

- Consider a dataset  $x_1, \ldots, x_n \sim F$
- Is the dataset from an Exp(λ) for some λ? I.e., is it F = Exp(λ)?
- We estimate  $\hat{\lambda} = 1/\bar{x}_n$
- We measure how close is the dataset to the distribution as:

$$t_{ks} = \sup_{a \in \mathbb{R}} |F_n(a) - F_{\hat{\lambda}}(a)|$$

where:

- $F_n(a)$  is the empirical cumulative distribution of  $x_1, \ldots, x_n$
- $F_{\hat{\lambda}}(a) = 1 e^{\hat{\lambda} a}$ , for  $a \ge 0$ , is the distribution function of  $Exp(\hat{\lambda})$
- ► t<sub>ks</sub> is called the Kolmogorov-Smirnov distance
- if  $F = Exp(\lambda)$  then both  $F_n \approx F$  and  $F_{\hat{\lambda}} \approx F$ , and then  $F_n \approx F_{\hat{\lambda}}$ , so that  $t_{ks}$  is small
- if  $F \neq Exp(\lambda)$  then  $F_n \approx F \neq Exp(\lambda) \approx F_{\hat{\lambda}}$ , so that  $t_{ks}$  is large

#### See R script

# Application: distribution fitting

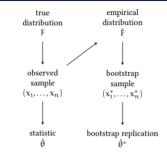
- For the software dataset from the textbook
  - $\hat{\lambda} = 0.0015$  and  $t_{ks} = 0.17$
- Is  $t_{ks} = 0.17$  expected or an extreme value?
- Let's study the distribution of the bootstrap estimator:

$${T_{ks}} = \mathop {\sup }\limits_{a \in \mathbb{R}} \left| {F_n^* (a) - F_{{{\hat \Lambda }^*}} (a)} 
ight|$$

where:

- $X_1^*, \ldots, X_n^* \sim \textit{Exp}(\hat{\lambda})$  is a bootstrap sample
- $F_n^*(a)$  is the empirical cumulative distribution of the bootstrap sample •  $\hat{\Lambda}^* = 1/\bar{X}_n^*$
- It turns out  $P(T_{ks} > 0.17) \approx 0$ , unlikely that Exp() is the right model

### See R script



### Hypothesis testing

- In the previous application, we tested how likely is Exp() for the given dataset
- In general, hypotheses testing consists of contrasting two conflicting theories (hypotheses) based on observed data
- Consider the German tank problem:
  - Military intelligence states that N = 350 tanks were produced
  - Alternative hypothesis:
    - N < 350 (one-tailed or one-sided test), or  $N \neq 350$  (two-tailed or two-sided test)
  - Observed serial tank id's: 61 19 56 24 16
- Statistical test: How likely is the observed data under the null hypothesis?
  - ▶ If it is NOT (sufficiently) likely, we reject the null hypothesis in favor of H1
  - ▶ If it is (sufficiently) likely, we cannot reject the null hypothesis
- Why 'we cannot reject the null hypothesis' and not instead 'we accept the null hypothesis'?
  - ▶ Other hypotheses, e.g., N = 349 or N = 351, could also not be rejected
  - We cannot say which of N = 349 or N = 350 or N = 351 is actually true

[H0 or null hypothesis]

[H1 hypothesis]

### Test statistic

TEST STATISTIC. Suppose the dataset is modeled as the realization of random variables  $X_1, X_2, \ldots, X_n$ . A *test statistic* is any sample statistic  $T = h(X_1, X_2, \ldots, X_n)$ , whose numerical value is used to decide whether we reject  $H_0$ .

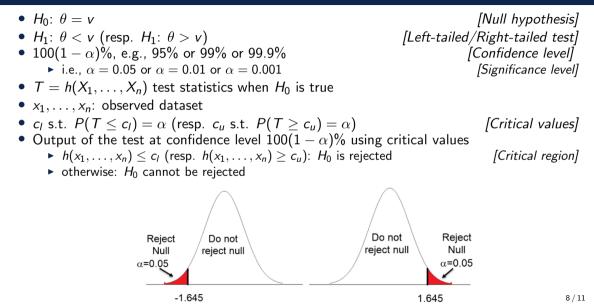
- In the German tank example:
  - $H_0: N = 350$
  - ► *H*<sub>1</sub> : *N* < 350
  - Observed serial tank id's: 61 19 56 24 16
- We use  $T = \max \{X_1, X_2, X_3, X_4, X_5\}$
- If  $H_0$  is true, i.e., N = 350, then  $E[T] = \frac{5}{6}(N+1) = \frac{5}{6}351 = 292.5$

Values in	Values in	Values against
favor of $H_1$	favor of $H_0$	both $H_0$ and $H_1$
5	292.5	1 350

• If H<sub>0</sub> is true, we have:

$$P(T \le 61) = P(\max\{X_1, X_2, X_3, X_4, X_5\} \le 61) = \frac{61}{350} \cdot \frac{60}{349} \dots \frac{57}{346} = 0.00014$$
very unlikely: either we are unfortunate, or  $H_0$  can be rejected

### Statistical test of hypothesis: one-tailed



## Statistical test of hypothesis: one-tailed

•  $H_0: \theta = v$ 

$$H_{1}: \theta < v \text{ (resp. } H_{1}: \theta > v \text{)} \qquad [Left-tailed/Right-tailed test]}$$

$$100(1 - \alpha)\%, \text{ e.g., } 95\% \text{ or } 99\% \text{ or } 99.9\% \text{ or } 99.9\% \text{ or } 99.9\% \text{ or } 90.01 \text{ or } \alpha = 0.001 \qquad [Confidence level]}$$

$$\bullet \text{ i.e., } \alpha = 0.05 \text{ or } \alpha = 0.01 \text{ or } \alpha = 0.001 \qquad [Significance level]}$$

$$T = h(X_{1}, \dots, X_{n}) \text{ test statistics when } H_{0} \text{ is true}$$

$$x_{1}, \dots, x_{n}: \text{ observed dataset}$$

$$p = P(T \le h(x_{1}, \dots, x_{n})) \text{ (resp. } p = P(T \ge h(x_{1}, \dots, x_{n}))) \qquad [p-value]$$

$$\bullet \text{ evidence against } H_{0} \text{ - the smaller the stronger evidence}$$

$$Output \text{ of the test at confidence level } 100(1 - \alpha)\% \text{ using } p\text{-values}$$

$$\bullet p \le \alpha: H_{0} \text{ is rejected}$$

$$\bullet \text{ otherwise: } H_{0} \text{ cannot be rejected}$$

p=0.009

α=0.05

[Null hypothesis]

## Statistical test of hypothesis: two-tailed

• 
$$H_0: \theta = v$$
 [Null hypothesis]  
•  $H_1: \theta \neq v$  [Two-tailed test]  
•  $100(1 - \alpha)\%$ , e.g., 95% or 99% or 99.9% [Confidence level]  
• i.e.,  $\alpha = 0.05$  or  $\alpha = 0.01$  or  $\alpha = 0.001$  [Significance level]  
•  $T = h(X_1, \dots, X_n)$  test statistics when  $H_0$  is true  
•  $x_1, \dots, x_n$ : observed dataset  
•  $c_l$  s.t.  $P(T \leq c_l) = \alpha/2$  and  $c_u$  s.t.  $P(T \geq c_u) = \alpha/2$  [Critical values]  
•  $h(x_1, \dots, x_n) \leq c_l$  or  $h(x_1, \dots, x_n) \geq c_u$ :  $H_0$  is rejected  
• otherwise:  $H_0$  cannot be rejected  
Reject  
Null  
 $\alpha/2 = 0.025$  Do not  
Reject Null  
 $\alpha/2 = 0.025$ 

1.96

-1.96

# Type I and Type II errors

		True state of nature	
		$H_0$ is true	$H_1$ is true
Our decision on the basis of the data	Reject $H_0$	Type I error	Correct decision
	Not reject $H_0$	Correct decision	Type II error

• Type I error: we falsely reject  $H_0$ 

[ $\alpha$ -risk, false positive rate]

 $[\beta$ -risk, false negative rate]

- ► E.g., convicting an innocent defendant
- ▶ we reject  $H_0$  when  $p < \alpha$ , so this error occur with probability  $100\alpha\%$
- $\blacktriangleright$  this error can be controlled by setting the significance level  $\alpha$  to the largest acceptable value
- how much is an acceptable value?
- A possible solution is to solely report the *p*-value, which conveys the maximum amount of information and permits decision makers to choose their own level
- Type II error: we falsely do not reject  $H_0$ 
  - E.g., acquitting a criminal
  - ▶  $1 \beta = P(\text{Reject}H_0 | H_1 \text{ is true})$  is called the *power* of the test