# **HYPOTHESIS TESTING • MTH107**

Class R FAQ

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# 1-Sample Z-Test

**H<sub>0</sub>:**  $\mu = \mu_0$  (where  $\mu_0$  = specific value)

Statistic:  $\overline{x}$  Test Statistic:  $Z = \frac{\overline{x} - \mu_0}{\frac{\sigma}{\sqrt{n}}}$  Conf. Region:  $\overline{x} + Z^* \frac{\sigma}{\sqrt{n}}$ 

**Assumptions:** 1)  $\sigma$  is known

2) n>30, n>15 and popn not strongly skewed, OR popn is normal

R: z.test()

# 1-Sample t-Test

**H<sub>0</sub>:**  $\mu = \mu_0$  (where  $\mu_0$  = specific value)

Statistic:  $\overline{x}$  Test Statistic:  $t=\frac{\overline{x}-\mu_0}{\frac{S}{\sqrt{n}}}$  Conf. Region:  $\overline{x}+t^*\frac{s}{\sqrt{n}}$  df: n-

**Assumptions:** 1)  $\sigma$  is UNknown,

2) n≥40, n≥15 & histogram not strongly skewed, OR histogram is normal

R: t.test(), hist()

# 2-Sample t-Test

 $H_0$ :  $\mu_1 = \mu_2$  Statistic:  $\overline{x}_1 - \overline{x}_2$ 

Conf. Region:  $(\bar{x}_1 - \bar{x}_2) + t^* \sqrt{s_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}$  df:  $n_1 + n_2 - 2$ 

**Assumptions:** 

- 1) Individuals in populations are independent
- 2) Variances are equal (use Levene's Test)
- 3) n₁+n₂≥40, n₁+n₂≥15 and both histograms are not strongly skewed, OR both histograms are normal

R: t.test(), levenesTest(), hist()

# **Choosing a Hypothesis Test**

1. If response variable is QUANTITATIVE, then GOTO 2; otherwise GOTO 5.

### **Quantitative Response**

- 2. If 1 POPULATION was sampled, then GOTO 3; otherwise GOTO 4.
- 3. If  $\sigma$  is KNOWN, then 1-Sample Z; otherwise 1-Sample t.
- If individuals in populations are INDEPENDENT, then 2-Sample t; otherwise, Paired t.

### **Categorical Response**

5. If 1 POPULATION was sampled, then Goodness-of-Fit; otherwise, Chi-Square.

# Making a Decision about H<sub>0</sub>

If the p-value  $< \alpha$ , then **REJECT H<sub>0</sub>**, otherwise **DNR H<sub>0</sub>**.

# **Chi-Square Test**

 $H_0$ : "Distribution of individuals into response levels is the same for all populations"  $H_a$ : "Distribution of individs into response levels is NOT the same for all populations"

Statistic: Observed frequency table

Test Statistic:  $\chi^2 = \sum \frac{(\text{Observed-Expected})^2}{\text{Expected}}$  df: (rows-1)(columns-1)

**Assumptions:** ≥5 in each cell of the expected table

R: xtabs(), matrix(), chisq.test(), percTable()

### **Goodness-of-Fit Test**

H<sub>0</sub>: "Distribution of individs into response levels follows the theoretical distribution"

**H**<sub>A</sub>: "Distribution of indivuals into response levels does NOT follow the theoretical distribution"

Statistic: Observed frequency table

Test Statistic:  $\chi^2 = \sum \frac{(\text{Observed-Expected})^2}{\text{Expected}}$  df: cells-1

**Assumptions:** ≥5 in each cell of the expected table

R: xtabs(), c(), chisq.test(), percTable(), chiGOF()

# 11 STEPS FOR ANY HYPOTHESIS TEST

- 1) State the rejection criterion  $(\alpha)$
- 2) State the null & alternative hypotheses and define the parameter(s)
- Determine which test to perform Explain!
- 4) Collect the data (address type of study and randomization)
- Check all necessary assumption(s)
- 6) Calculate the appropriate statistic(s)
- 7) Calculate the appropriate test statistic
- 8) Calculate the p-value
- 9) State your rejection decision
- 10) Summarize your findings in terms of the problem
- 11) If rejected  $H_0$ , compute a 100(1- $\alpha$ )% confidence region for parameter