One-Sample Hypothesis Tests for Multivariate Mean Vectors

ST 558: Multivariate Analytics

Module 3

Lecture 2

We will first quickly review the univariate case to recall the vocabulary and philosophy of hypothesis testing:

- Let $X_1, X_2, ..., X_n$ be an iid sample from a population with population mean μ and population variance σ^2 .
- Question/Goal: Test hypotheses regarding the value of the (unknown) mean parameter μ .

Hypotheses to be tested:

• Null hypothesis:

$$H_0: \mu = \mu_0$$

for some specified value μ_0

• We compare this null hypothesis to an *Alternative hypothesis*. We can consider several forms for the alternative hypothesis:

$$H_A: \mu > \mu_0$$
 OR

$$H_A: \mu < \mu_0$$
 OR

$$H_A$$
: $\mu \neq \mu_0$

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We will focus on the third of these possible alternatives, the *two-sided alternative hypothesis*.

Example: We have measured body temperature on n = 12 healthy subjects.

| Subject | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|------------|------|------|------|------|------|------|------|------|------|------|------|------|
| Body Temp. | 94.3 | 95.2 | 98.2 | 96.8 | 99.3 | 98.5 | 99.4 | 97.6 | 99.2 | 98.3 | 97.9 | 98.0 |

We would like to use this data to perform test of the null hypothesis that the population mean (average) body temperature is 98.6° F.

$$H_0$$
: $\mu = 98.6$

$$H_A$$
: $\mu \neq 98.6$

The result of a hypothesis test is one of two possible decisions/outcomes:

- Reject the null hypothesis H_0 : conclude that the data do not support the null hypothesis
- Fail to reject the null hypothesis H_0 : conclude that the data could possibly support the null hypothesis

We decide between these two outcomes using a *test* statistic, which is a function of the observed data.

The value of the test statistic determines the decision.

There are two different errors we could make as a result of the hypothesis testing decision:

- Type I error: Reject H_0 when H_0 is in fact true
- Type II error: Fail to reject H_0 when H_0 is not true.

| | Truth | | | | | | |
|----------------------|------------------|------------------|--|--|--|--|--|
| Decision | H_0 | NOT H_0 | | | | | |
| Reject H_0 | Type I Error | Correct Decision | | | | | |
| Fail to Reject H_0 | Correct Decision | Type II Error | | | | | |

Hypothesis tests are designed to control the probability of making a Type I error:

$$\alpha = P(\text{Type I error})$$

= $P(\text{Reject } H_0 \text{ when } H_0 \text{ is true})$
= $P_{H_0}(\text{Reject } H_0)$

This probability α is called the *significance level* of the test.

The probability of making a Type II error depends on the <u>true value</u> of the parameter being tested.

Often instead of talking about the probability of a Type II error, we talk about the *power* of a test procedure at a specific value μ_A of the parameter tested

Power(
$$\mu_A$$
) = $P(\text{Reject } H_0 \text{ when } \mu_A \text{ is true})$
= $P_{\mu_A}(\text{Reject } H_0)$

When comparing tests with the same significance level, we prefer a test with *higher power*, since that means the test will make the correct decision more often.

Univariate Hypothesis Test for Population Mean

Recall the *t*-test:

- Setting: $X_1, ..., X_n$ iid from a population with unknown mean μ (and unknown variance σ^2)
- Null hypothesis:

$$H_0: \mu = \mu_0$$

Alternative hypothesis:

$$H_A$$
: $\mu \neq \mu_0$

Test statistic:

$$t(\mu_0) = \frac{\sqrt{n}(\bar{X} - \mu_0)}{s}$$

• Decision:

Reject
$$H_0$$
 at level α for $|t(\mu_0)| > t_{(n-1)}(\alpha/2)$

Univariate Hypothesis Test for Population Mean

Recall the *t*-test:

- Setting: λ Throughout this course, we will only be considering tests that **do** mean μ (not assume that the population variance (or covariance matrix)
- Null hype is known (the textbook does cover the case where the variance is known, however).
- Alternati
 This is because in practice, the variance/covariance is pretty much never known if the population mean is not known.
- Test statistic:

$$t(\mu_0) = \frac{\sqrt{n}(\bar{X} - \mu_0)}{s}$$

• Decision: Reject H_0 at level α for $|t(\mu_0)| > t_{(n-1)}(\alpha/2)$

Univariate Hypothesis Test for Population Mean

Recall the *t*-test

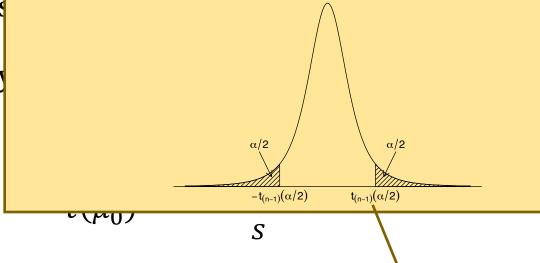
• Setting: X_1 , ... mean μ (and

Null hypothes

Alternative hypersulation

• Test statistic:

The <u>critical value</u> $t_{(n-1)}(\alpha/2)$ is the upper $\alpha/2$ quantile of a *t*-distribution with (n-1) degrees of freedom:



• Decision:

Reject H_0 at level α for $|t(\mu_0)| > t_{(n-1)}(\alpha/2)$

Example: We have measured body temperature on n = 12 healthy subjects.

| Subject | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|------------|------|------|------|------|------|------|------|------|------|------|------|------|
| Body Temp. | 94.3 | 95.2 | 98.2 | 96.8 | 99.3 | 98.5 | 99.4 | 97.6 | 99.2 | 98.3 | 97.9 | 98.0 |

We would like to use this data to perform a level $\alpha = 0.05$ test of the null hypothesis that the population mean (average) body temperature is 98.6° F.

$$H_0$$
: $\mu = 98.6$

$$H_A$$
: $\mu \neq 98.6$

Example: We have measured body temperature on n = 12 healthy subjects.

| | Subject | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|----|-----------|------|------|------|------|------|------|------|------|------|------|------|------|
| Во | ody Temp. | 94.3 | 95.2 | 98.2 | 96.8 | 99.3 | 98.5 | 99.4 | 97.6 | 99.2 | 98.3 | 97.9 | 98.0 |

• Test statistic:

$$t(98.6) = \frac{\sqrt{12}(97.725 - 98.6)}{\sqrt{2.518}} = -1.910$$

Critical value:

$$t_{(11)}(0.025) = qt(1-0.025, 11) = 2.201$$

• Decision: Since $|t(98.6)| < t_{(11)}(0.025)$, we **fail to reject** the null hypothesis that the population mean is 98.6.

Example: We have measured body temperature on n = 12 healthy subjects.

| Subject | 1 | Sample mean | 5 | 6 | 7 | 8 | Sample variance | 12 |
|------------|------|--------------------|------|------|------|------|-----------------|-----|
| Body Temp. | 94.3 | $\bar{X} = 97.735$ | 99.3 | 98.5 | 99.4 | 97.6 | $s^2 = 2.518$ | 8.0 |

• Test statistic:

$$t(98.6) = \frac{\sqrt{12}(97.725 - 98.6)}{\sqrt{2.518}} = -1.910$$

Critical value:

$$t_{(11)}(0.025) = qt(1-0.025, 11) = 2.201$$

• Decision: Since $|t(98.6)| < t_{(11)}(0.025)$, we **fail to reject** the null hypothesis that the population mean is 98.6.

Now, with multivariate data $X_1, X_2, ..., X_n$, we want to test hypotheses regarding a *mean vector*

$$\mathbf{\mu} = \left[\mu_1, \mu_2, \dots, \mu_p\right]^T$$

• Null hypothesis:

$$H_0$$
: $\mu = \mu_0$

• Alternative hypothesis:

$$H_A$$
: $\mu \neq \mu_0$

How can this be done?

Example: We have measured body temperature and pulse on *n* = 12 healthy subjects.

| Subject | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|------------|------|------|------|------|------|------|------|------|------|------|------|------|
| Body Temp. | 94.3 | 95.2 | 98.2 | 96.8 | 99.3 | 98.5 | 99.4 | 97.6 | 99.2 | 98.3 | 97.9 | 98.0 |
| Pulse | 72 | 75 | 76 | 78 | 73 | 79 | 77 | 70 | 79 | 72 | 87 | 76 |

We would like to use this data to perform a level $\alpha = 0.05$ test of the null hypothesis that the population mean vector is (98.6° F, 77 bpm).

$$H_0$$
: $\mu = (98.6, 77)$

$$H_A$$
: $\mu \neq (98.6, 77)$

One idea for testing a hypothesis H_0 : $\mu = \mu_0$:

- Just test each variable's mean separately: Using only the observations for variable j, perform a level α test of H_{0j} : $\mu_j = \mu_{0j}$, where
 - $\circ \mu_j$ is the true (unknown) population mean of the *j*th variable
 - $\circ \mu_{0j}$ is the *j*th element of the hypothesized mean vector μ_0
- Reject H_0 if **any** of the H_{0j} are rejected.

Will this work? Will this approach provide a level α test of H_0 : $\mu = \mu_0$?

Remember, our goal is to achieve

$$P_{H_0}(\text{Reject } H_0: \boldsymbol{\mu} = \boldsymbol{\mu}_0) = \alpha$$

If we test each individual variable at level α , then we will *inflate our Type I error probability*, because we are giving ourselves multiple opportunities to make a Type I error: this is a *multiple testing* scenario.

Therefore, this approach *will not* provide a level α test of H_0 : $\mu = \mu_0$.

We could correct for the *multiple testing* issue using a *Bonferroni* correction:

Test each individual variable's mean at level

$$\alpha^* = \frac{\alpha}{p}$$

(where p is the number of variables measured, as usual).

Then we would have:

$$P_{H_0}(\text{Reject } H_0: \mathbf{\mu} = \mathbf{\mu}_0) = P_{H_0}(\text{Reject at least one of } H_{0j})$$

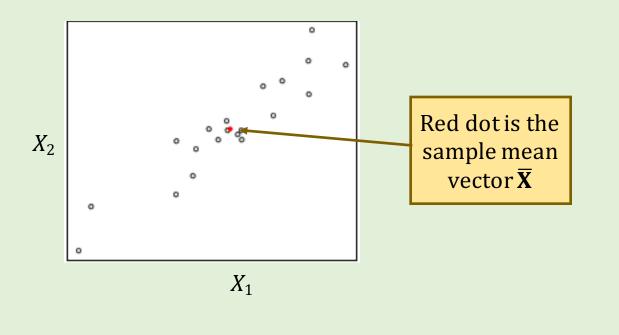
$$\leq \sum_{j=1}^p P_{H_{0j}}(\text{Reject } H_{0j}) = \sum_{j=1}^p \frac{\alpha}{p} = \alpha$$

So the probability of a Type I error would be controlled: guaranteed to be less than or equal to the desired level α .

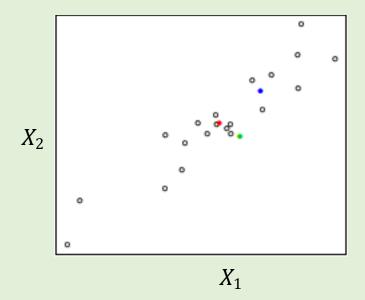
However, in many cases performing individual tests using a Bonferroni correction and combining the results will not be a desirable solution:

- This procedure is too <u>conservative</u> for most settings: it rejects a true null hypothesis with probability strictly less than the desired significance level α .
- Conservatism generally implies a lack of power: this test will not reject the null in many settings where the null is in fact false.

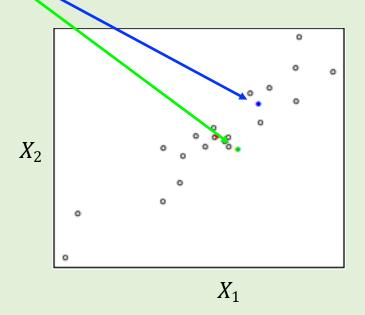
Suppose we have the bivariate data represented in this scatterplot:



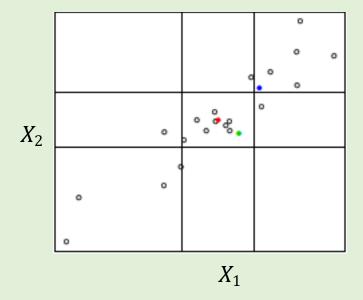
Which of these two points (blue or green) do you think is more plausible as the population mean vector?



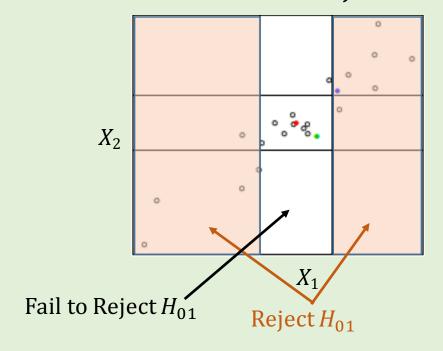
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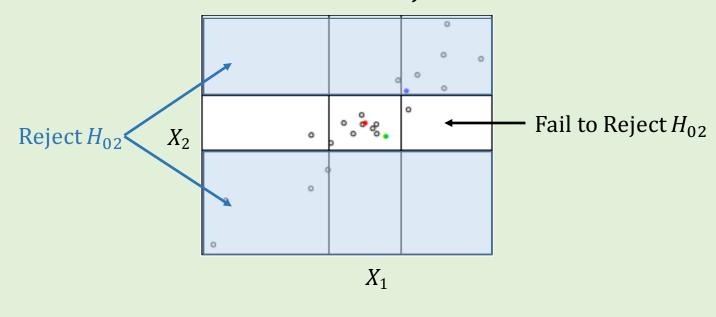
If we performed univariate tests of each variable's mean separately, the black lines indicate the range of population mean values that would NOT be rejected:



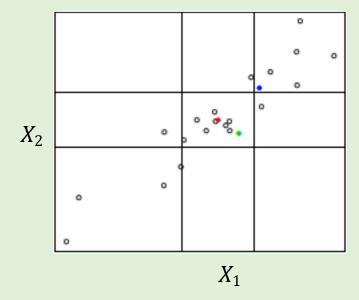
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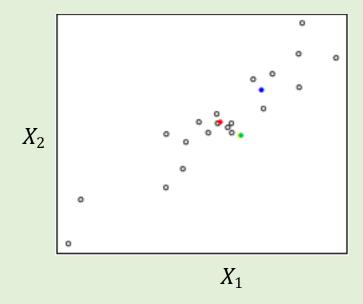
If we performed univariate tests of each variable's mean separately, the black lines indicate the range of population mean values that would NOT be rejected:



Therefore, if we test each variable's mean separately, we would *reject* the blue point, but *fail to reject* the green point as plausible population mean vectors.



However, we might convincingly argue that actually the blue point looks more reasonable than the green point: it is more centered within the data cloud.



The separate univariate tests do not take into account the relationship *between* the variables, but <u>that relationship</u> <u>can give us useful information</u>.

