STAT 339: Statistical Theory

Frequentist Hypothesis Testing

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Elements of a Statistical Test

Introduction

Objective: Make inferences about unknown population parameters

• Based on information contained in sample data

We can estimate the parameter using, for example, point or interval estimation.

• We can also test specific claims about the parameter.

A (Frequentist) statistical hypothesis test involves four components:

- 1. Null hypothesis, H_0
- 2. Alternative hypothesis, H_A
- 3. Test statistic
- 4. Rejection region

My Awesome Function

I wrote an **R function** called **generateCat** that returns either "Moose" or "Cannoli". You don't know anything else about it.

I claim that this function returns "Moose" 50% of the time.

Do you trust me?1

```
generateCat(n = 20)
```

► Output

[1] No, you definitely don't trust me. You believe the "Moose" proportion, p, is higher than 50%.

Null and Alternative Hypothesis

Because you **do not believe me** when I claim that the "Moose" proportion is 50%, you might seek to support the research hypothesis that the "Moose" proportion is higher than 50%.

• This research hypothesis is commonly known as the alternative hypothesis.

Support for the alternative hypothesis is obtained by showing (using sample data as evidence) that the null hypothesis is false.

- The null hypothesis is the converse of the alternative hypothesis:
 - $\cdot H_0: p = 0.5$
 - $\circ \ H_A: p > 0.5$

Our research objective is to show whether the data support rejecting the null hypothesis in favor of the alternative.

Test Statistic

In order to decide between the null and alternative hypothesis, we need data.

• The **test statistic** (similar to an estimator) is a function of the **sample data**, Y_1, \ldots, Y_n .

```
generateCat(n = 20)

### n Moose_count Moose_prop
### 1 20     11     0.55
```

A test statistic for this test is Y=11. This is the number of times $\operatorname{generateCat}$ returned "Moose".

• Our statistical decision will be based on this test statistic.

Rejection Region

The **rejection region** (RR) specifies the values of the test statistic for which the null hypothesis is to be rejected in favor of the alternative hypothesis.

- We reject the null hypothesis, in favor of the alternative, if a particular sample yields a test statistic that falls in the RR.
- If the value of the test statistic does not fall into the RR, we fail to reject the null hypothesis.

In our example, larger values of Y would lead us to reject H_0 .

ullet For example, one RR might be the set of all values of $Y\geq 16$:

$$RR = \{y : y \ge 16\}$$

But is this a good rejection region? It's clear that larger values of Y, say $\mathbf{y} \geq \mathbf{k}$ are more contradictory to $H_0: p=0.5$, but how should we choose k?

Errors in Hypothesis Testing

For any rejection region, two types of erors can be made in reaching a decision for our test:

- Type I Error: Reject H_0 when H_0 is true.
 - The probability of a type I error is denoted by α , which is also called the level of the test.
- Type II Error: Fail to reject H_0 when H_A is true.
 - \circ The probability of a type II error is denoted by β .

Clearly, we don't want to make incorrect decisions!

• Because α and β measure the risks associated with the two possible incorrect decisions we might reach, they can help us find a "good" rejection region.

Type I Error

In our example, I ran generate $\mathsf{Cat}\,n=20$ times.

We wish to test

$$H_0: p = 0.5$$
 versus $H_A: p > 0.5$

where p is the $\it true$ proportion of times this function is supposed to return "Moose".

- The test statistic is Y, the number of sampled times that generateCat returned "Moose".
- Let's find lpha, using $RR=\{y\geq 16\}$.

```
lpha = P(	ext{type I error}) = P(	ext{reject } H_0 	ext{ when } H_0 	ext{ is true}) \ = P(	ext{value of test statistic is in RR when } H_0 	ext{ is true}) \ = P(Y \ge 16 \mid p = 0.5)
```

```
1 - pbinom(15, size = 20, prob = 0.5) # P(Y >= 16 | p = 0.5)
```

Type II Error

Using $RR = \{y \geq 16\}$, is our test equally good in protecting us from type II error?

• That is, do we have low risk of concluding that the "Moose" proportion is equal to 50% when in fact it is greater?

```
eta = P(	ext{type II error}) = P(	ext{fail to reject } H_0 	ext{ when } H_A 	ext{ is true}) \ = P(	ext{value of test statistic is NOT in RR when } H_A 	ext{ is true}) \ = P(Y < 16 \mid p = 0.6)
```

ullet Note: We used p=0.6 as a particular value of p that satisfies H_A .

```
pbinom(15, size = 20, p = 0.6)
```

[1] 0.949048

Type I and II Error

When using $RR = \{y \geq 16\}$...

- $\alpha = 0.006$ (low risk of type I error)
- $\beta = 0.949$ when p = 0.6 (high risk of type II error)

We can balance lpha and eta by enlarging the RR. Suppose instead we used:

$$RR = \{y \ge 13\}$$

Partier solution: Increase the sample size!!!

Common Large-Sample Tests

Introduction

Goal: Test a set of hypotheses concerning a parameter θ , based on a random sample Y_1, Y_2, \ldots, Y_n .

Setting: Hypothesis testing based on an estimator $\hat{\theta}$ that has (approximately) a normal sampling distribution with mean θ and standard error $\sigma_{\hat{\theta}}$.

• That is,

$$\hat{ heta} \sim N(heta, \sigma_{\hat{ heta}})$$

Our large-sample tests will be based on the large-sample estimators from earlier this semester:

- ullet $ar{Y}$
- ullet \hat{p}
- $\bar{Y}_1 \bar{Y}_2$
- $oldsymbol{\hat{p}}_1 \hat{p}_2$

Large-Sample Estimators

(From WMS, page 397)

$\begin{array}{c} {\rm Target} \\ {\rm Parameter} \\ \theta \end{array}$	Sample Size(s)	Point Estimator $\hat{\theta}$	$E(\hat{ heta})$	Standard Error $\sigma_{\hat{ heta}}$
μ	n	\overline{Y}	μ	$\frac{\sigma}{\sqrt{n}}$
p	n	$\hat{p} = \frac{Y}{n}$	p	$\sqrt{\frac{pq}{n}}$
$\mu_1 - \mu_2$	n_1 and n_2	$\overline{Y}_1 - \overline{Y}_2$	$\mu_1 - \mu_2$	$\sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}^{*\dagger}$
$p_1 - p_2$	n_1 and n_2	$\hat{p}_1 - \hat{p}_2$	$p_1 - p_2$	$\sqrt{\frac{p_1q_1}{n_1} + \frac{p_2q_2}{n_2}}^{\dagger}$

Setting Up The Test

Suppose we wish to test

$$H_0: \theta = \theta_0 \quad \text{versus} \quad H_A: \theta > \theta_0$$

where θ_0 is a specific value of θ .

- If in reality $heta> heta_0$, then $\hat{ heta}$ is more likely to be large.
 - \circ In this case, larger values of $\hat{ heta}$ favor rejection of H_0 .

The components of our statistical hypothesis test are as follows:

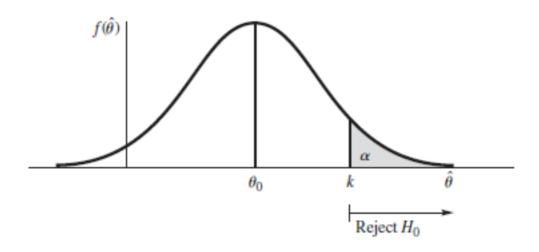
- 1. Null Hypothesis: $H_0: \theta = \theta_0$
- 2. Alternative Hypothesis: $H_A: heta > heta_0$
- 3. Test Statistic: $\hat{\theta}$
- 4. Rejection Region: $RR = \{\hat{ heta} > k\}$ for some choice of k

The Rejection Region

To determine k in the **rejection region**, we'll fix the **type 1 error** probability α (the **level** of the test) and choose k accordingly.

• $\alpha = P(ext{type I error}) = P(ext{reject } H_0 ext{ when } H_0 ext{ is true}) = 0.05 ext{ (usually)}$

WMS Figure 10.3 (page 496)



The Rejection Region

Remember, θ is fixed but unknown.

• But when H_0 is true, $heta= heta_0$, and by the Central Limit Theorem,

$$\hat{ heta} \sim N(heta_0, \sigma_{\hat{ heta}}) \iff Z = rac{\hat{ heta} - heta_0}{\sigma_{\hat{ heta}}} \sim N(0, 1)$$

Therefore, an equivalent rejection region would be \$\$RR=\left{\hat{\theta}:Z=\frac{\hat{\theta}-\theta{0}} {\sigma{\hat{\theta}}}>z{\alpha}\right}, \$\$ where `(P(Z>z{\alpha})=\alpha)`

Large-Sample (right tailed) Test

A large-sample test, with level α , is as follows:

- 1. Null Hypothesis: $H_0: heta= heta_0$
- 2. Alternative Hypothesis: $H_A: heta > heta_0$
- 3. Test Statistic: $Z=rac{\hat{ heta}- heta_0}{\sigma_{\hat{ heta}}}$
- 4. Rejection Region: $RR = \{z > z_{lpha}\}$ (upper-tail rejection region)

Example

The level of our test is $\alpha = 0.05$.

• This means $RR = \{z > z_{0.05}\}$:

```
qnorm(0.05, lower.tail = FALSE)
```

[1] 1.644854

My Awesome Function

I wrote an **R function** called **generateCat** that returns either "Moose" or "Cannoli". You don't know anything else about it.

I claim that this function returns "Moose" 50% of the time.

You don't trust me and definitely think it's higher. Test this hypothesis using an $\alpha=0.05$ level.

```
generateCat(n = 40)
```

```
## n Moose_count Moose_prop
## 1 40 24 0.6
```

- $H_0: p = 0.5$
- $H_A: p > 0.5$

From our sample, n=40 and $\hat{p}=0.6$.

The Null Distribution

Under H_0 , $p_0=0.5$, and by the Central Limit Theorem,

$$Z = rac{\hat{p} - p_0}{\sqrt{rac{p_0(1-p_0)}{n}}} \sim N(0,1)$$

for sufficiently large n.

- Note: While we could've used $\sqrt{\hat{p}(1-\hat{p})/n}$ to approximate the standard error of \hat{p} , we use p_0 because we are considering the distribution of Z under H_0 .
- $RR = \{z > 1.645\}$
- $Z=rac{0.6-0.5}{\sqrt{rac{0.5(1-0.5)}{40}}}=1.265$

Decision: Fail to reject $H_0!$ At the lpha=0.05 level, the evidence does not support your decision to not trust me...

- Does this mean I'm telling the truth?! 😇
- Only time will tell... We need to calculate the probability of type II error, β .

Two-Tailed Tests

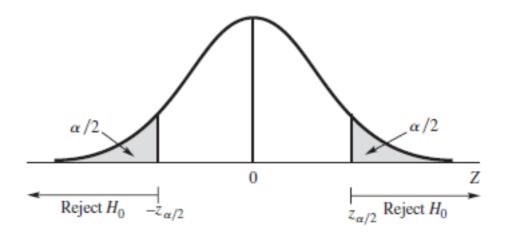
Suppose we wanted to test an alternative hypothesis of the form: $heta
eq heta_0$.

ullet That is, we're testing $H_0: heta = heta_0$ versus $H_A: heta
eq heta_0$

Our test statistic would remain the same!

$$Z = rac{\hat{ heta} - heta_0}{\sigma_{\hat{ heta}}}$$

• The one thing that would change is the **rejection region**. That's because, in this setting, we'll reject H_0 if $\hat{\theta}$ is (a) much smaller or (b) much larger than θ_0 .



Common Grounds

I want to test whether the **mean** number of people in line at **Common Grounds** on a typical morning after STAT 339, μ , is **different from** 5 people.

 For all I know, the mean could be < 5 or > 5. Considering I can't even get the name right, I don't know much else about Common Grounds...



I collect some DATA and obtain the following:

- n=30 (yes, I checked the Common Grounds line for 30 mornings after STAT 339)
- ullet $ar{y}=6$ people in line, with standard deviation s=2

Do the data present sufficient evidence to suggest that the mean number of people in line at Common Grounds after STAT 339 is different from 5 people? Use $\alpha=0.01$.

Summary

Large-Sample lpha-Level Hypothesis Tests

- 1. Null Hypothesis: H_0 : $\theta = \theta_0$
- 2. Alternative Hypothesis:
 - $\circ H_A: \theta > \theta_0$ (upper-tail alternative)
 - $\circ~H_A$: $heta < heta_0$ (lower-tail alternative)
 - $\circ H_A$: $heta
 eq heta_0$ (two-tailed alternative)
- 3. Test Statistic: $Z=rac{\hat{ heta}- heta_0}{\sigma_{\hat{ heta}}}$
- 4. Rejection Region:
 - $\circ \ \{z>z_{lpha}\}$ (upper-tail RR)
 - $\circ \{z < z_{lpha}\}$ (lower-tail RR)
 - $\circ~\{|z|>z_{lpha/2}\}$ (two-tailed RR)

Type II Error Probabilities

Type II Error

Recall: Type II Error occurs when you fail to reject the null, when the alternative is true.

For a test with $H_0: \theta = \theta_0$ versus $H_A: \theta > \theta_0$, we can calculate type II error probabilities only for specific values of θ in H_A .

ullet For some specific alternative, $heta= heta_A$, the probability of type II error, eta, is

$$eta = P(\hat{ heta} ext{ is not in RR when } H_A ext{ is true}) \ = P(\hat{ heta} \le k \mid heta = heta_A) \ = P\left(rac{\hat{ heta} - heta_A}{\sigma_{\hat{ heta}}} \le rac{k - heta_A}{\sigma_{\hat{ heta}}} \mid heta = heta_A
ight),$$

By the Central Limit Theorem, $Z=(\hat{ heta}- heta_A)/\sigma_{\hat{ heta}}\sim N(0,1)$, so we can calculate

$$eta = P\left(Z \leq rac{k - heta_A}{\sigma_{\hat{ heta}}} \mid heta = heta_A
ight)$$

by finding a corresponding area under the N(0,1) curve, using ${\tt pnorm}(\,).$

Type II Error Probability

In My Awesome Function example, we had n=40 and $\hat{p}=0.6$. Let's find eta for a test of

$$H_0: p = 0.5$$
 versus $H_A: p = 0.65$.

ullet From the example, the **rejection region** for an lpha=0.05 level test was

$$z = rac{\hat{p} - p_0}{\sqrt{rac{p_0(1-p_0)}{n}}} > 1.645 \iff \hat{p} > p_0 + 1.645 \sqrt{rac{p_0(1-p_0)}{n}}.$$

ullet Plugging in $p_0=0.5$ and n=40, we have the rejection region $\{\hat{p}>0.63\}.$

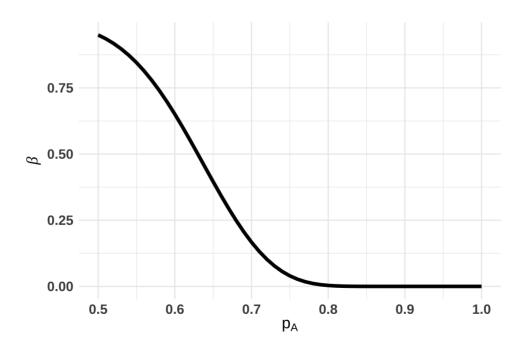
Therefore, by definition, the type II error probability is

$$eta = P(\hat{ heta} ext{ is not in RR when } H_A ext{ is true}) = P(\hat{p} \le 0.63 \mid p = 0.65)$$
 $= P\left(Z \le rac{0.63 - 0.65}{\sqrt{rac{0.65(1 - 0.65)}{40}}}
ight)$
 $= P(Z \le -0.265) = \boxed{0.396}$

Understanding β

While we can only calculate β for a fixed p_A , we can visualize how β depends on the distance between p_A and p_0 (0.5).

• If p_A is far from p_0 , then the true value of p is a bit easier to detect, hence β is much smaller.



Back to Piéchart Emporium (yet again)

Piéchart Emporium claims that they pull in an average of 1000 dollars in daily revenue, but you definitely think it's lower. To check this, you could test

$$H_0: \mu = 1000$$
 versus $H_A: \mu < 1000$

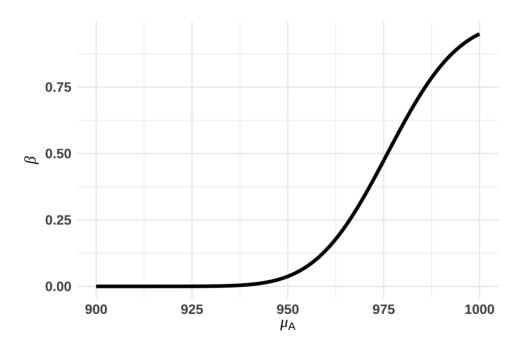
at the $\alpha=0.05$ level, using the **rejection region** $\{z<-1.645\}$ and the following observed data:

- n=30 days
- $\bar{y}=950$ dollars
- s=80 dollars

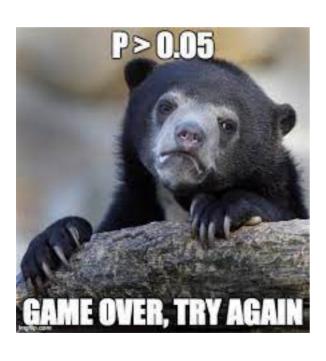
In groups of 3, calculate β for one of the following three specific values of μ under H_A :

- $1.\,H_A:\mu=980$
- 2. $H_A: \mu = 960$
- 3. $H_A: \mu = 940$

Understanding β



p-values



my p value: < 0.05 the null hypothesis:



What is a p-value?

- **p-value**: The probability of observing a test statistic at least as extreme as the one you observed, if the null hypothesis is true.
- In other words, this is the smallest level of significance, α , for which the observed data indicate that the null hypothesis should be rejected.
- The smaller the p-value becomes, the stronger the evidence that the null hypothesis should be rejected.

Typically, we use p-values in the following way to reach decisions in a hypothesis test:

- $p \le \alpha$: Reject the null hypothesis
- $p > \alpha$: Fail to reject the null hypothesis

But be careful with this line of thinking! There is more to science than an arbitrary, binary decision making process!

My Awesome Function

In the generateCat example, we tested

$$H_0: p = 0.5$$
 versus $H_A: p > 0.5$

using n=40 and $\hat{p}=0.6$.

• Using lpha=0.05, our rejection region was $RR=\{z>1.645\}$, and our observed test statistic was

$$Z = rac{0.6 - 0.5}{\sqrt{rac{0.5(1 - 0.5)}{40}}} = 1.265.$$

Under H_0 , $Z \sim N(0,1)$, so

p-value =
$$P(Z \ge 1.265 \mid H_0 \text{ is true}) = 0.103$$

1-pnorm(1.265)

[1] 0.1029357

Common Grounds

I want to test whether the **mean** number of people in line at **Common Grounds** on a typical morning after STAT 339, μ , is **different from** 5 people.

- $H_0: \mu = 5$
- $H_A: \mu \neq 5$



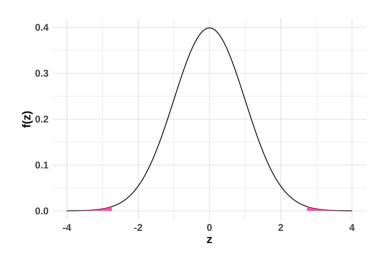
Data:
$$n = 30, \bar{y} = 6, s = 2$$

• Though rather than rejecting H_0 for only large values of \bar{y} , we might also reject H_0 for small values of \bar{y} . When calculating the p-value, we need to account for this!

Common Grounds

Our test statistic for this test was

$$Z = \frac{6-5}{2/\sqrt{30}} = 2.74.$$



The pink regions in the plot represent the p-value for a two-tailed test.

In R:

[1] 0.006143918

Decision: p < 0.05, so reject the null hypothesis.

Relationship with Confidence Intervals

(In my opinion) Confidence intervals are more informative and less confusing than p-values.

 Not that p-values are useless; the definition just lends itself well to misinterpretation!

There's also a duality that exists between large-sample CIs and two-tailed hypothesis tests:

- Do not reject $H_0: \theta = \theta_0$ at the α level if θ_0 lies inside a $100(1-\alpha)\%$ confidence interval for θ .
- Reject H_0 if θ_0 lies outside the interval.

Example

95% CI for the mean number of people in line at Common Grounds on a typical morning after STAT 339, μ :

$$6\pm1.96 imesrac{2}{\sqrt{30}}=(5.28,6.72)$$

Small-Sample Hypothesis Testing

(for means)

Introduction

The large-sample hypothesis testing procedures rely on the condition that the sample size is large enough that

$$Z = rac{\hat{ heta} - heta_0}{\sigma_{\hat{ heta}}} \sim (approx) \ N(0,1).$$

A common procedure that we can use with **small samples** relies on using the t-distribution as the sampling distribution of the **test statistic**.

 This procedure, the t-test, is only appropriate for samples from Normal populations!

The t-test

Setting

- Y_1, Y_2, \ldots, Y_n denote a random sample of size n
- The sample comes from a Normal distribution with unknown mean μ and unknown variance σ^2 .

Procedure

- Let \bar{Y} and S denote the sample mean and sample standard deviation, respectively.
- ullet Under the **null hypothesis**, $H_0: \mu = \mu_0$, the test statistic

$$T=rac{ar{Y}-\mu_0}{S/\sqrt{n}}\sim t(n-1).$$

- For $H_A: \mu > \mu_0$, the rejection region is $RR = \{t > t_{lpha}\}.$
 - \circ That is, any value of T such that $P(T>t_{lpha})=lpha$, where t_{lpha} is the upper lpha-quantile of the t(n-1) distribution.

Florida Lobsters



WMS 10.75

A declaration by the Bahamian government that prohibits U.S. lobsterers from fishing on the Bahamian portion of the continental shelf was expected to dramatically reduce the landings (in pounds per lobster trap).

According to records, the prior mean landings per trap was 30.31 pounds.

A random sample of **20 lobster traps** since the Bahamian fishing restriction went into effect gave the following results (in pounds):

17.4, 18.9, 39.6, 34.4, 19.6, 33.7, 37.2, 43.4, 41.7, 27.5, 24.1, 39.6, 12.2, 25.5, 22.1, 29.3, 21.1, 23.8, 43.2, 24.4

Do these landings provide evidence that the mean landings per trap has decreased? Test using lpha=0.05

Comparing Groups

We can also use the t-distribution to **compare the means** of two Normal populations that possess equal variances.

Setting

• Independent samples from Normal distributions with $\sigma_1^2=\sigma_2^2$.

Hypotheses

- $H_0: \mu_1 \mu_2 = D_0$, where D_0 is usually 0.
- ullet $H_A: \mu_1-\mu_2>D_0$ (or < or eq depending on which alternative is appropriate)

Comparing Groups

We can also use the t-distribution to **compare the means** of two Normal populations that possess equal variances.

Test Statistic

$$ullet$$
 $T=rac{(ar{Y}_1-ar{Y}_2)-D_0}{S_p\sqrt{rac{1}{n_1}+rac{1}{n_2}}}$, where

$$S_p^2 = rac{(n_1-1)S_1^2 + (n_2-1)S_2^2}{n_1 + n_2 - 2}$$

Rejection Region

• For an upper-tail alternative, $RR=\{t>t_{lpha}\}$, where t_{lpha} is the upper lpha-quantile of the $t(n_1+n_2-2)$ distribution.

Is the sequel always better?

I watched **Shrek** recently, because Netflix decided it was going to pull the movie at the end of April 2022.

• Not sure yet what this means for **Shrek 2**, and as of this writing I haven't watched it in a while. But let's compare them anyways!



Suppose random samples of $n_1=n_2=10$ critic reviews from Metacritic for each film yielded the ratings below. Check if the sequel is better at $\alpha=0.01$.

- \bullet Shrek: 89.62, 82.03, 79.65, 88.20, 82.13, 86.06, 90.33, 89.44, 86.27, 79.82
- Shrek 2: 77.05, 73.93, 76.27, 70.89, 79.65, 64.49, 67.39, 74.15, 71.50, 86.16

Connecting to Confidence Intervals

The connection between two-tailed **t-tests** and small-sample confidence intervals is similar to the duality between large-sample confidence intervals and tests!

- Do not reject $H_0: \mu = \mu_0$ if μ_0 lies inside $ar y \pm t_{lpha/2} imes rac{S}{\sqrt n}$
- Reject $H_0: \mu = \mu_0$ if μ_0 lies outside $ar y \pm t_{lpha/2} imes rac{S}{\sqrt{n}}$
- (Ditto for CIs for a difference between means)
 - \circ Reject $H_0: \mu_1 \mu_2 = D_0$ if D_0 lies outside the CI.

Example:

Calculate a 99% confidence interval for the difference in average Metacritic ratings between Shrek and Shrek 2.

t-tests in R

Because the **t-test** is a common hypothesis testing procedure, it is supported by the **base R t.test()** function.

For example, we can repeat the *lobster* example in R using the following code:

```
### One Sample t-test
### data: lobster_data
### t = -0.64678, df = 19, p-value = 0.2628
### alternative hypothesis: true mean is less than 30.31
### 95 percent confidence interval:
### -Inf 32.61098
### sample estimates:
### mean of x
### 28.935
```

Testing Hypotheses Concerning Variances

Introduction

Recall: If $Y_1,Y_2,\ldots,Y_n\sim N(\mu,\sigma^2)$, where both μ and σ^2 are unknown, then

$$rac{(n-1)S^2}{\sigma^2} \sim \chi^2(n-1).$$

We can use this fact to test $H_0:\sigma^2=\sigma_0^2$ for some fixed value σ_0^2 .

If $H_0:\sigma^2=\sigma_0^2$ is true, then

$$\chi^2 \equiv rac{(n-1)S^2}{\sigma_0^2} \sim \chi^2(n-1),$$

and we can use χ^2 as our test statistic.

• For an lpha-test of $H_0:\sigma^2=\sigma_0^2$ versus $H_A:\sigma^2>\sigma_0^2$,

$$RR = \{\chi^2 > \chi^2_{lpha}\}$$

Example

(WMS 10.79)

The manufacturer of a machine to package soap power claimed that their machine could load cartons at a given weight with a range of no more than 0.4 ounce. The mean and variance of a sample of n=8 three-pound boxes were found to equal 3.1 and 0.018, respectively.

Test the hypothesis that the variance of the population of weight measurements is $\sigma^2=0.01$ against the alternative that $\sigma^2>0.01$. Use $\alpha=0.05$.

Comparing Groups

Sometimes rather than examining one group's variance, we might want to compare the variances of two Normal distributions.

Usually to determine whether they are equal.

From Exam 2:

• If independent samples of size n_1 and n_2 are taken from two normally distributed populations with variances σ_1^2 and σ_2^2 , respectively, then

$$F = rac{S_1^2/\sigma_1^2}{S_2^2/\sigma_2^2} \sim F(df_1 = n_1 - 1, df_2 = n_2 - 1)$$

Suppose we're testing $H_0:\sigma_1^2=\sigma_2^2$ versus $H_A:\sigma_1^2>\sigma_2^2$. Under H_0 , $\sigma_1^2=\sigma_2^2$ and

$$F = rac{S_1^2/\sigma_1^2}{S_2^2/\sigma_2^2} = rac{S_1^2}{S_2^2} \sim F(df_1 = n_1 - 1, df_2 = n_2 - 1)$$

is a test statistic!

90s Road Trip

Because the test statistic, F, follows an $F(n_1-1,n_2-1)$ distribution under the null hypothesis, a **rejection region** for an α -level test comparing variances is

$$RR = \{F > F_{\alpha}\},$$

where F_{α} is the lpha-quantile of the $F(n_1-1,n_2-1)$ distribution.

Example

	90s Road Trip	No Music
Sample size, n	100	100
Mean, $ar{y}$	10.75	12.75
Standard deviation, \boldsymbol{s}	4	5

Using the data from Exam 2, is there sufficient evidence to conclude that the variance of *objects memorized* for the "No Music" group is *larger* than the variance for the "90s Road Trip" group?