22S:105 Statistical Methods and Computing

More on t-tests

Lecture 16 Mar. 23, 2011

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```
data normtemp;
infile 'normtemp.dat';
input temp gender heart;
run;

proc means n mean stddev clm alpha = .05;
var temp;
run;
```

Variable	N	Mean	Std Dev	Lower 95.0% CLM	Upper 95.0
temp	130	98.249	0.733	98.122	98.376

Note that:

- The 95% confidence interval for μ does not contain 98.6.
- The p-value is less than .05, so we can reject the null hypothesis.

SAS for one-sample t-tests

• SAS automatically does a two-sided test

$$H_0: \mu = \mu_0$$

 $H_a: \mu \neq \mu_0$

Example: Using the "normtemp.dat" data on body temperatures measured on 130 healthy adults, we will test the hypotheses

$$H_0: \mu = 98.6$$

 $H_a: \mu \neq 98.6$

at the .05 significance level.

Example 2: We will use the same dataset to test a hypothesis regarding heart rates, namely:

$$H_0: \mu = 73$$

 $H_a: \mu \neq 73$

at the .05 significance level.

Analysis Variable : HEART

N	Mean	Std Dev	Lower 95.0% CLM	Upper 95.0% CL
130	73.7615385	7.0620767	72.5360699	74.9870071

Note that:

- The 95% confidence interval for μ does contain 73.
- The p-value is greater than .05, so we cannot reject the null hypothesis.

One-sample t-tests using proc univariate

```
data normtemp ;
infile '/group/ftp/pub/kcowles/datasets/normtemp.dat' ;
input temp gender heart ;
format gender sexfmt. ;
run ;

proc univariate mu0 = 98.6 data = normtemp ;
var temp ;
run ;
```

The UNIVARIATE Procedure
Variable: temp

Tests for Location: Mu0=98.6

Test	-St	tatistic-	p Value	
Student's t	t	-5.45482	Pr > t	<.0001
Sign	M	-21	Pr >= M	0.0002
Signed Rank	S	-1963	Pr >= S	<.0001

Two-sample t-tests

So far we have talked about drawing inference about a single population mean μ based on data contained in one sample drawn from that population.

Now we will consider procedures for comparing two *different* population means.

There are different procedures depending on whether the samples are

- paired
- independent

Paired samples

- We are interested in the unknown population means μ_1 and μ_2 of two different populations.
- In our sample, each observation drawn from the first population is matched up with an observation drawn from the second population.

• *self-pairing*: two measurements are taken on each subject

Example:

- systolic blood pressure (sbp) upon entry into a clinical study
- -sbp after 1 month on treatment

The population means of interest are

- $-\mu_1$ = mean sbp of untreated patients of this type
- $-\mu_2$ = mean sbp of patients of this type after 1 month of treatment with the study regimen
- The question of interest is whether the treatment lowers blood pressure, i.e. is $\mu_2 < \mu_1$?

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• matched pairs: investigator matches each subject in one treatment group with one subject in another treatment group so that members of a pair are as alike as possible

The population means of interest are

- $-\mu_1$ = mean response (say sbp at 1 month) of patients receiving treatment 1
- $-\mu_2$ = mean response of patients receiving treatment 2
- The question of interest is whether $\mu_1 = \mu_2$

come OC users. This subgroup will be the study sample.

- Measure the sbp of the study sample at the follow-up visit.
- We will compare the baseline and follow-up sbps of the women in the study sample.

Paired t-test

To carry out the hypothesis test of interest, we apply one-sample procedures to the *differences* between values measured on members of each pair.

Example:

- We are interested in whether the use of oral contraceptive (OC) drugs affects the level of systolic blood pressure (sbp) in women.
- We identify a group of nonpregnant, premenopausal women aged 16-49 from a prepaid health plan who are not currently OC users and measure their sbp, which we will refer to as baseline sbp.
- We rescreen these women 1 year later to ascertain a subgroup who have remained non-pregnant throughout the year and have be-

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We will do a two-sided test, because we do not know in advance whether to expect μ_1 (mean sbp in OC users) to be higher or lower than μ_2 (mean sbp in non-users).

$$H_0: \mu_1 = \mu_2$$

 $H_a: \mu_1 \neq \mu_2$

or equivalently:

$$H_0: \mu_1 - \mu_2 = 0$$

 $H_a: \mu_1 - \mu_2 \neq 0$

or equivalently:

$$H_0: \delta = 0$$
$$H_a: \delta \neq 0$$

where δ denotes $\mu_1 - \mu_2$.

We will use the *observed differences* between the before and after values observed on each woman as our data to to carry out the hypothesis test regarding δ at the .05 significance level. data sbpoc ;
infile '/group/ftp/pub/kcowles/datasets/sbpoc.dat' ;
input sbpnooc sbpoc ;
diff = sbpoc - sbpnooc ;
run ;

proc print ;
run ;

OBS	SBPNOOC	SBPOC	DIFF
1	115	128	13
2	112	115	3
3	107	106	-1
4	119	128	9
5	115	122	7
6	138	145	7
7	126	132	6
8	105	109	4
9	104	102	-2
10	115	117	2

We will compute the sample mean of the d_i s

$$\bar{d} = \frac{\sum_{i=1}^{n} d_i}{n}$$

and the sample standard deviation of the d_i s

$$s_d = \sqrt{\frac{\Sigma_i^n (d_i - \bar{d})^2}{n - 1}}$$

proc means data = sbpoc ;
var diff ;
run ;

Analysis Variable : DIFF

N	Mean	Std Dev	Minimum	Maximum
10	4.800	4.5655716	-2.0000	13.0000

Then the t statistic is

$$t = \frac{\bar{d} - \delta_0}{s_d / \sqrt{n}}$$

From our data,

$$\begin{split} \bar{d} &= 4.80 \\ s_d &= 4.566 \\ t &= \frac{4.80}{4.566/\sqrt{10}} \\ &= 3.32 \end{split}$$

Using Table A, we see that the value that cuts off the upper .025 area under a t distribution with 9 degrees of freedom is 2.262.

Because 3.32 > 2.262 (our result is more extreme than the required cutoff), we can reject the null hypothesis at the .05 level.

We could use SAS to find the exact p-value, which is 0.0089.

Note that the one-sample t-test in proc univariate by default tests the null hypothesis that $\mu=0$.

```
proc univariate data = sbpoc ;
var diff ;
run ;
```

The UNIVARIATE Procedure Variable: temp

Tests for Location: Mu0=0

Test	t -Statistic-		p Value	
Student's t	t	3.324651	Pr > t	0.0089
Sign	M	3	Pr >= M	0.1094
Signed Rank	S	24	Pr >= S	0.0117