## STA 250/MTH 342 Intro to Mathematical Statistics Lab Session 9 / March 30, 2015 / Handout

In this lab session we review and implement some hypothesis tests.

See: https://stat.duke.edu/courses/Spring15/sta250/labs/ for links to source code and data. Submit lab solutions via email to: sta250@stat.duke.edu. Any plots should be included in postscript form as attachments. The email subject must be "STA250 ..." with "..." replaced by your name.

1: One-sample *t*-test. Suppose we have independent data  $X_1, \dots, X_n$  from No $(\mu, \sigma^2)$ , and the variance  $\sigma^2$  is unknown. We do the following test,

$$H_0: \mu = \mu_0$$
 vs.  $H_1: \mu \neq \mu_0$ .

In the lecture notes we defined

$$T = \frac{\sqrt{n}(X_n - \mu_0)}{s} \sim t_{n-1},$$

and the corresponding rejection region

$$\mathscr{R}(C) = \{ |T| > C \}.$$

Now we to the test in **R**, and we set  $\mu_0 = 0$ .

```
> base <- "https://stat.duke.edu/courses/Spring15/sta250/labs/lab9";
> download.file(paste(base,"data1.Rdata",sep="/"),"data1.Rdata","wget");
> load("data1.Rdata");
> data1;
  \begin{bmatrix} 1 \end{bmatrix} \quad 0.7787533 \quad 0.9435898 \quad -0.1453734 \quad -0.6031512 \quad 0.8058303 \quad -2.4178814 \\ \end{bmatrix} 
 [7] 0.8380311 -0.3673249 -0.3779325 -1.2226983 -0.1546096 2.4170250
[13] 0.4929351 0.6918879 -0.9982076 1.0864768 -1.1788555 -0.4409982
[19] 1.4905478 0.3973025
> conf.level <- 0.95;
> alter <- c("two.sided", "less", "greater")[1];</pre>
> t.test(data1, alternative = alter, conf.level = conf.level);
         One Sample t-test
data: data1
t = 0.4103, df = 19, p-value = 0.6861
alternative hypothesis: true mean is not equal to 0
95 percent confidence interval:
 -0.4173234 0.6208581
sample estimates:
mean of x
0.1017674
```

In the test report generated by  $\mathbf{R}$ , "t" is our statistic T. The variable "df" is the degree of freedom. In this case n = 20 so df = 19.

The "p-value" is the *p*-value of the statistic, defined as the smallest  $\alpha$  level at which observing **X** will lead to a rejection of the test. So, if **R** reports a "p-value" smaller than the test level  $\alpha$ , we should

reject the null hypothesis. In this example, we do not reject the null hypothesis since the p-value is greater than  $\alpha = 0.05$ .

The "conf.level" is the confidence level  $1 - \alpha$ , where  $\alpha$  is the level of the test. The confidence interval is the one for  $\mu$  if we forget about the hypothesis testing task.

Confidence Interval = 
$$\left[ \bar{X}_n + \frac{F_{t_{19}}^{-1}(\frac{\alpha}{2})}{\sqrt{n}}, \ \bar{X}_n + \frac{F_{t_{19}}^{-1}(1-\frac{\alpha}{2})}{\sqrt{n}} \right].$$

Therefore one may also use the t.test function to find confidence interval.

This test is illustrated in Figure 1.

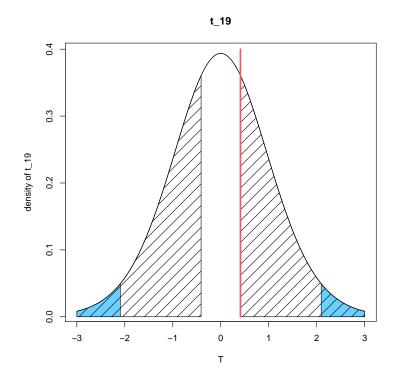


Figure 1: The two-sided *t*-test with level  $\alpha = 0.05$ . The blue part marks the rejection region of area  $\alpha$ . The red line marks *T*. The area of the shaded part is the *p*-value.

For plotting the power function, recall that for any  $\mu_1 \neq \mu_0$ , and any  $0 < \sigma < \infty$ ,

$$\pi(\mu_1,\sigma) = \Pr(\mathscr{R}(F_{t_{19}}^{-1}\left(1-\frac{\alpha}{2}\right))|\mu_1,\sigma) = F_{t_{19}(\psi)}\left(F_{t_{19}}^{-1}\left(\frac{\alpha}{2}\right)\right) + 1 - F_{t_{19}(\psi)}\left(F_{t_{19}}^{-1}\left(1-\frac{\alpha}{2}\right)\right),$$

where  $\psi = (\mu_1 - \mu_0)\sqrt{n}/\sigma$ . Below is the code for plotting this power function, and the plot is in Figure 2.

```
alpha <- 0.05
C <- qt(1-alpha/2, df = 19)
power.fun <- function(mu,sigma){
  return(pt(-C, df = 19, ncp = (mu-0)*sqrt(20)/sigma) + 1 -
    pt(C, df = 19, ncp = (mu-0)*sqrt(20)/sigma))
}
```

```
mu <- seq(-3,3,length.out = 30)
sigma <- seq(0.1, 5, length.out = 30)
z <- outer(mu, sigma, Vectorize(power.fun))
surface3d(mu, sigma, z)
# require(rgl)
# persp(mu, sigma, z, theta = -50, phi=30)</pre>
```

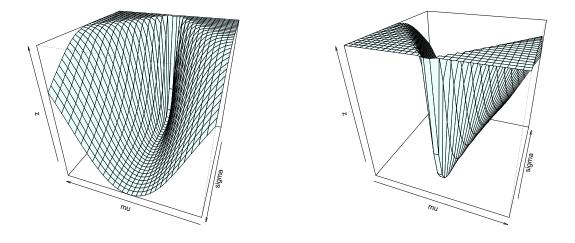


Figure 2: The power function of the above test.  $\mu_1$  runs from -3 to 3 and  $\sigma$  runs from 0.1 to 5.

**TASK 1** With same data please do *t*-test with the one-sided alternative

$$H_0: \mu \le \mu_0$$
 vs.  $H_1: \mu > \mu_0$ ,

where please set  $\mu_0 = 0$ . This could be done use the "alternative = "greater" argument in the "t.test" function. Please also plot its power function. You may need page 21, in lecture note 17 for reference.

2: Two-sample *t*-test, with common variance. Now suppose we have

$$X_1, \cdots, X_n \sim \mathsf{No}(\mu_1, \sigma^2),$$
  
$$Y_1, \cdots, Y_m \sim \mathsf{No}(\mu_2, \sigma^2).$$

Consider the following test

$$H_0: \mu_1 = \mu_2$$
 vs.  $H_1: \mu_1 \neq \mu_2$ .

Recall the test statistic

$$T = \frac{X - Y}{\sqrt{\frac{(n-1)s_X^2 + (m-1)s_Y^2}{n+m-2}} \cdot \sqrt{\frac{1}{m} + \frac{1}{n}}} \sim t_{n+m-2}.$$

Recall the rejection region for level  $\alpha$ ,

 $\mathscr{R}(C) = \{ |T| > C \}, \quad C = F_{t_{n+m-2}}^{-1} \left( 1 - \frac{\alpha}{2} \right).$ 

Let's do the test in **R**. We use  $\alpha = 0.05$ .

```
> download.file(paste(base, "data2.Rdata", sep="/"), "data2.Rdata", "wget");
> load("data2.Rdata");
> conf.level <- 0.95;
> alter <- c("two.sided", "less", "greater")[1];
> t.test(X, Y, alternative = alter, conf.level = conf.level, var.equal = T);
        Two Sample t-test
data: X and Y
t = -3.2534, df = 53, p-value = 0.001987
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
        -2.0691000 -0.4908667
sample estimates:
    mean of x mean of y
0.0869087 1.3668920
```

Since "p-value = 0.001987", which is less than  $\alpha$ , the null hypothesis is rejected.

**TASK 2** Use the above data, test the hypothesis

 $H_0: \mu_1 \le \mu_2$  vs.  $H_1: \mu_1 > \mu_2$ .

## 3: Welch's approximate *t*-test. Now suppose we have

$$X_1, \cdots, X_n \sim \mathsf{No}(\mu_1, \sigma_1^2)$$
$$Y_1, \cdots, Y_m \sim \mathsf{No}(\mu_2, \sigma_2^2).$$

Consider the following test

$$H_0: \mu_1 = \mu_2$$
 vs.  $H_1: \mu_1 \neq \mu_2$ 

Recall the Welch's approximate *t*-test, where a *t*-statistic (remember that this is just an approximation, this is strictly speaking not a *t*-statistic!)

$$T_w = \frac{X - Y}{\sqrt{\frac{s_X^2}{n} + \frac{s_Y^2}{m}}}$$

The exact sampling distribution of  $T_w$  under  $H_0$  is not known but it is very close to a t distribution with the following degrees of freedom,

df = 
$$\frac{(s_X^2/n + s_Y^2/m)^2}{\frac{(s_X^2/n)^2}{n-1} + \frac{(s_Y^2/m)^2}{m-1}}$$
.

We now do the test, just using the above data.

```
> load("data2.Rdata");
> conf.level <- 0.95;
> alter <- c("two.sided", "less", "greater")[1];
> t.test(X, Y, alternative = alter, conf.level = conf.level, var.equal = F);
        Welch Two Sample t-test
data: X and Y
t = -3.3755, df = 52.949, p-value = 0.001386
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
        -2.040572 -0.519395
sample estimates:
mean of x mean of y
0.0869087 1.3668920
```

We see that the degree of freedom is no longer an integer. Since the p-value is small, the null hypothesis is rejected.

**TASK 3** Use the above data, test the hypothesis

 $H_0: \mu_1 \le \mu_2$  vs.  $H_1: \mu_1 > \mu_2$ .

4: Analyzing paired data with one-sample *t*-test. Suppose  $X_1, \dots, X_n$  are the numbers of hours of sleep that *n* individuals get on Day 1. Suppose that after some treatment, we measure the hours of sleep they get on Day 2:  $Y_1, \dots, Y_n$ . Recall that in the lecture notes we treat this kind of data as paired samples, where for each *i*,  $X_i$  may not be independent with  $Y_i$ . We consider the hypothesis

```
H_0: \mu_1 = \mu_2 vs. H_1: \mu_1 \neq \mu_2.
```

```
> download.file(paste(base,"data1.Rdata",sep="/"),"data1.Rdata","wget");
> load("data1.Rdata");
> conf.level <- 0.95;</pre>
> alter <- c("two.sided", "less", "greater")[1];</pre>
> t.test(X, Y, alternative = alter, conf.level = conf.level,
    var.equal = T, paired = T);
+
        Paired t-test
data: X and Y
t = -0.7304, df = 22, p-value = 0.4729
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -1.3397268 0.6418685
sample estimates:
mean of the differences
             -0.3489292
```

Since the *p*-value is greater than  $\alpha$ , we do not reject the null hypothesis.

**TASK 4** Using the above data, do the test

 $H_0: \mu_1 \ge \mu_2$  vs.  $H_1: \mu_1 < \mu_2$ .

6

 ${\sim}{\sim}{\rm END}{\sim}{\sim}$