# STA305/1004 - Class 7

January 30, 2017

### Today's Class

- ► Case study on power poses study: study replication and power
- Sample size and power in studies with two proportions
- Calculating power via simulation
- ▶ Introduction to causal inference



Professor Amy Cuddy

#### Cuddy's study methods:

- Randomly assigned 42 participants to the high-power pose or the low-power-pose condition.
- Participants believed that the study was about the science of physiological recordings and was focused on how placement of electrocardiography electrodes above and below the heart could influence data collection.
- Participants' bodies were posed by an experimenter into high-power or low-power poses. Each participant held two poses for 1 min each.
- Participants' risk taking was measured with a gambling task; feelings of power were measured with self-reports.
- Saliva samples, which were used to test cortisol and testosterone levels, were taken before and approximately 17 min after the power-pose manipulation.

(Carney, Cuddy, Yap, 2010)

#### Cuddy's study results:

As hypothesized, high-power poses caused an increase in testosterone compared with low-power poses, which caused a decrease in testosterone, F(1, 39) = 4.29, p < .05; r = .34. Also as hypothesized, high-power poses caused a decrease in cortisol compared with low-power poses, which caused an increase in cortisol, F(1, 38) = 7.45, p < .02; r = .43

- ▶ The study was replicated by Ranehill et al. (2015)
- An initial power analysis based on the effect sizes in Carney et al. (power = 0.8,  $\alpha$  = .05) indicated that a sample size of 100 participants would be suitable.

```
library(pwr)
pwr.t.test(d=0.6,power = 0.8)
```

Two-sample t test power calculation

```
n = 44.58579
d = 0.6
sig.level = 0.05
power = 0.8
alternative = two.sided
```

NOTE: n is number in \*each\* group

- ▶ Ranehill et al. study used a sample of 200 participants to increase reliability.
- ▶ This study found none of the significant differences found in Cuddy's study.
- ▶ The replication study obtained very precise estimates of the effects.
- What happened?

- ▶ Sampling theory predicts that the variation between samples is proportional to  $\frac{1}{\sqrt{n}}$ .
- In small samples we can expect variability.
- Many researchers often expect that these samples will be more similar than sampling theory predicts.

# Study replication

Suppose that you have run an experiment on 20 subjects, and have obtained a significant result from a two-sided z-test ( $H_0: \mu=0$  vs. $H_1: \mu\neq 0$ ) which confirms your theory ( $z=2.23,\ p<0.05,$  two-tailed). The researcher is planning to run the same experiment on an additional 10 subjects. What is the probability that the results will be significant at the 5% level by a one-tailed test ( $H_1: \mu>0$ ), seperately for this group?

STUDY 1: Ho: 
$$\mu = 0$$
 Us  $44$ :  $\mu \neq 0$   
 $Z = 2.23 = \frac{X-0}{0/\sqrt{20}} = 2.23 \frac{0}{\sqrt{120}}$ 

$$\frac{x-0}{5/50} > 1.645 \Rightarrow x > 1.645 = 5$$

Power = 
$$P\left(\frac{X > 1.645}{X > 1.645}\right)$$
 To , when  $M = 2.23$   $O/\sqrt{520}$ 

$$= P\left(\frac{X - M}{\sqrt{500}} > \frac{1.645}{\sqrt{500}}\right) \frac{1.645}{\sqrt{500}}$$

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$$=P(2>0.068)=0.47$$

- In many clinical trials, the primary endpoint is dichotomous, for example, whether a patient has responded to the treatment, or whether a patient has experienced toxicity.
- ▶ Consider a two-arm randomized trial with binary outcomes. Let  $p_1$  denote the response rate of the experimental drug,  $p_2$  as that of the standard drug, and the difference is  $\theta = p_1^- p_2$ .

Let  $Y_{ik}$  be the binary outcome for subject i in arm k; that is,

$$Y_{ik} = \left\{ egin{array}{ll} 1 & ext{with probability } p_k \ 0 & ext{with probability } 1-p_k, \end{array} 
ight.$$

for  $i = 1, ..., n_k$  and k = 1, 2. The sum of indendent and identically distributed Bernoulli random variables has a binomial distribution,

$$\sum_{i=1}^{n_k} Y_{ik} \sim Bin(n_k, p_k), \ k = 1, 2.$$

(Yin, pg. 173-174)

The sample proportion for group k is

$$\hat{p}_k = \bar{Y}_k = (1/n_k) \sum_{i=1}^{n_k} Y_{ik}, \ k = 1, 2,$$

and 
$$E\left(\bar{Y}_k\right)=p_k$$
 and  $Var\left(\bar{Y}_k\right)=rac{p_k(1-p_k)}{n_k}$ .

The goal of the clinical trial is to determine if there is a difference between the two groups using a binary endpoint. That is we want to test  $H_0: \theta = 0$  versus  $H_0: \theta \neq 0$ .

The test statistic (assuming that  $H_0$  is true) is:

$$T = rac{\hat{
ho}_1 - \hat{
ho}_2}{\sqrt{
ho_1(1-
ho_1)/n_1 + 
ho_2(1-
ho_2)/n_2}} \sim N(0,1),$$

The test rejects at level  $\alpha$  if and only if

$$|T| \geq z_{\alpha/2}$$
.

Using the same argument as the case with continuous endpoints and ignoring terms smaller than  $\alpha/2$  we can solve for  $\beta$ 

$$eta pprox \Phi \left( z_{lpha/2} - rac{| heta_1|}{\sqrt{
ho_1(1-
ho_1)/n_1 + 
ho_2(1-
ho_2)/n_2}} 
ight).$$

Using this formula to solve for sample size. If  $n_1 = r \cdot n_2$  then

$$n_2 = rac{\left(z_{lpha/2} + z_{eta}
ight)^2}{ heta^2} \left( p_1 (1-p_1)/r + p_2 (1-p_2) 
ight).$$

- The built-in R function power.prop.test() can be used to calculate sample size or power.
- ▶ For example suppose that the standard treatment for a disease has a response rate of 20%, and an experimental treatment is anticipated to have a response rate of 28%.
- ▶ The researchers want both arms to have an equal number of subjects. How many patients should be enrolled if the study will conduct a two-sided test at the 5% level with 80% power?

```
power.prop.test(p1 = 0.2,p2 = 0.25,power = 0.8)
```

 ${\tt Two-sample\ comparison\ of\ proportions\ power\ calculation}$ 

```
n = 1093.739
p1 = 0.2
p2 = 0.25
sig.level = 0.05
power = 0.8
alternative = two.sided
```

NOTE: n is number in \*each\* group

- If the test statistic and distribution of the test statistic are known then the power of the test can be calculated via simulation.
- ► Consider a two-sample t-test with 30 subjects per group and the standard deviation of the clinical outcome is known to be 1.
- ▶ What is the power of the test  $H_0: \mu_1 \mu_2 = 0$  versus  $H_0: \mu_1 \mu_2 = 0.5$ , at the 5% significance level?
- The power is the proportion of times that the test correctly rejects the null hypothesis in repeated sampling.

We can simulate a single study using the rnorm() command. Let's assume that  $n_1=n_2=30, \mu_1=3.5, \mu_2=3, \sigma=1, \alpha=0.05$ .

```
set.seed(2301)
t.test(rnorm(30,mean=3.5,sd=1),rnorm(30,mean=3,sd=1),var.equal = T)
```

Two Sample t-test

```
data: rnorm(30, mean = 3.5, sd = 1) and rnorm(30, mean = 3, sd = 1)
t = 2.1462, df = 58, p-value = 0.03605
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
    0.03458122   0.99248595
sample estimates:
mean of x mean of y
    3.339362   2.825828
```

Should you reject  $H_0$ ?

- Suppose that 10 studies are simulated.
- ▶ What proportion of these 10 studies will reject the null hypothesis at the 5% level?
- ▶ To investigate how many times the two-sample t-test will reject at the 5% level the replicate() command will be used to generate 10 studies and calculate the p-value in each study.
- ▶ It will still be assumed that  $n_1 = n_2 = 30, \mu_1 = 3.5, \mu_2 = 3, \sigma = 1, \alpha = 0.05.$

[1] 0.03604893 0.15477655 0.01777959 0.40851999 0.34580930 0.11131007 [7] 0.14788381 0.00317709 0.09452230 0.39173723

```
*power is the number of times the test rejects at the 5\% level sum(pvals \le 0.05)/10
```

[1] 0.3

But, since we only simulated 10 studies the estimate of power will have a large standard error. So let's try simulating 10,000 studies so that we can obtain a more precise estimate of power.

[1] 0.4881

This is much closer to the theoretical power obtained from power.t.test().

```
power.t.test(n = 30,delta = 0.5,sd = 1,sig.level = 0.05)
```

Two-sample t test power calculation

n = 30
delta = 0.5
sd = 1
sig.level = 0.05
power = 0.477841
alternative = two.sided

NOTE: n is number in \*each\* group

- ► The built-in R functions power.t.test() and power.prop.test() don't have an option for calculating power where the there is unequal allocation of subjects between groups.
- ► These built-in functions don't have an option to investigate power if other assumptions don't hold (e.g., normality).
- One option is to simulate power for the scenarios that are of interest. Another option is to write your own function using the formula derived above.

- Suppose the standard treatment for a disease has a response rate of 20%, and an experimental treatment is anticipated to have a response rate of 28%.
- ▶ The researchers want both arms to have an equal number of subjects.
- ► A power calculation above revealed that the study will require 1094 patients for 80% power.
- ▶ What would happen to the power if the researchers put 1500 patients in the experimental arm and 500 patients in the control arm?

- ► The number of subjects in the experimental arm that have a positive response to treatment will be an observation from a Bin(1500, 0.28).
- ► The number of subjects that have a positive response to the standard treatment will be an observation from a Bin(500, 0.2).
- We can obtain simulated responses from these distributions using the rbinom() command in R.

```
set.seed(2301)
rbinom(1,1500,0.28)

[1] 403
rbinom(1,500,0.20)
```

[1] 89

prop 1 prop 2 0.2686667 0.1780000

▶ The p-value for this simulated study can be obtained using prop.test().

```
correction

data: c(rbinom(1, 1500, 0.28), rbinom(1, 500, 0.2)) out of c(1500, 500
X-squared = 16.62, df = 1, p-value = 4.568e-05
alternative hypothesis: two.sided
95 percent confidence interval:
    0.05032654 0.13100680
sample estimates:
```

2-sample test for equality of proportions without continuity

- ▶ A power simulation repeats this process a large number of times.
- In the example below we simulate 10,000 hypothetical studies to calculate power.

[1] 0.6231

If the researchers decide to have a 3:1 allocation ratio of patients in the treatment to control arm then the power will be \_\_\_\_\_?