#### Lecture 10

### 10.1 Finishing up last example

- Consider this example where we left-off from last class:
  - \* In a Zogby Poll conducted with 1,203 likely voters nationwide between Sep 24-28, 2023, Donald Trump led Joe Biden, 52.5 percent to 47.5 percent, among those expressing a preference.
    - \* This is a tracking poll. In the previous three-day window of the poll (Sep 21-23), Trump led Biden 55.6 to 44.4 percent (N=1,203).
    - \* According to the poll, Trump's lead shrunk by about six points in three days. How confident are we that this change is not due to sampling error?
    - \* Set it up:
    - \* The parameter we seek is now  $p_1 p_2$ , where  $p_1$  = Trump's true support in the first poll (Sep 21-23) and  $p_2$  = Trump's true support in the second poll.
    - \* The polls may be considered two binomial experiments in which  $Y_1$  is the number of "successes" (here, the # favoring Trump) in the first poll, (no ideological agenda) and  $Y_2$  is the number of of such "successes" in the second poll.
    - \* An intuitive estimator for this quantity would be  $\hat{p}_1 \hat{p}_2$ , where the p-hats are the proportions of respondents favoring Trump in the two polls. Is it an unbiased estimator for  $p_1 p_2$ ?

$$E(\widehat{p}_1 - \widehat{p}_2) = E(\widehat{p}_1) - E(\widehat{p}_2)$$

$$= E\left(\frac{Y_1}{n_1}\right) - E\left(\frac{Y_2}{n_2}\right) [\widehat{p}_1 \text{ and } \widehat{p}_2 \text{ are functions of the RVs } Y_1, Y_2]$$

$$= \frac{1}{n_1} E(Y_1) - \frac{1}{n_2} E(Y_2)$$

$$= \frac{1}{n_1} n_1 p_1 - \frac{1}{n_2} n_2 p_2 [E(Y) = np \text{ if } Y \text{ is distributed binomial}]$$

$$= p_1 - p_2.$$

- \* Our next step is to say how precise  $\hat{p}_1 \hat{p}_2$  tends to be as an estimator of  $p_1 p_2$ .
- \* We do this by figuring out what the estimator's standard error is. It's

$$\sqrt{VAR(\widehat{p}_1 - \widehat{p}_2)} = \sqrt{VAR(\widehat{p}_1) + VAR(\widehat{p}_2)} \text{ [assume samples drawn independently]}$$

$$= \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$$

\* We make the substitution

$$(\widehat{p}_1 - \widehat{p}_2) \pm z_{\alpha/2} \sqrt{\frac{\widehat{p}_1(1-\widehat{p}_1)}{n_1} + \frac{\widehat{p}_2(1-\widehat{p}_2)}{n_2}}$$

\* Plugging in, we have

$$(55.6 - 52.5) \pm z_{\alpha/2} \sqrt{\frac{(55.6)(100 - 55.6)}{1,203} + \frac{(52.5)(100 - 52.5)}{1,203}}$$
  
 $3.1 \pm z_{\alpha/2}(2.031).$ 

\* Do you recall how we find  $z_{\alpha/2}$ ? We type qnorm( $\frac{\alpha}{2}$ ), substituting our chosen  $\alpha$ . You'll remember that  $z_{\alpha/2}$  associated with an  $\alpha=.05$  is  $z_{.025}=-1.96$ . So our 95% CI is:

$$3.1 \pm 1.96(2.031) = 3.1 \pm 3.98$$
, or  $[-.9, 7.1]$ .

- \* We are 95% confident that the true change between the two polls was between -.9 and 7.1 percentage points.
- Note that this CI includes zero. So another interpretation of this CI is that we are not 95% confident that there was zero change between the two polls. And this, of course, is what we really wanted to know: was there truly any movement between Oct 21-23 and Oct 24-26?
- Now, does the 90% confidence interval about our point estimate include zero?
  - Let's see: our alpha is .10.
  - typing qnorm(.05) gives us -1.64. So our 90% CI is:

$$3.1 \pm 1.64(2.031) = 3.1 \pm 3.33$$
, or [-.23, 6.43].

- Still no cigar. At what level of confidence would we be satisfied that there was movement between the two surveys?
- Think: we wish to find some  $\alpha^*$  such that the lower bound of the  $100 * (1 \alpha)$  CI is greater than zero. That is, find some  $\alpha^*$  meeting this criterion:

$$\alpha^* : 3.1 - z_{\alpha^*/2}(2.031) > 0.$$

• To do this, manipulate the expression

$$-z_{\alpha^*/2}(2.031) > -3.1$$

$$z_{\alpha^*/2} < \frac{3.1}{2.031}$$

$$z_{\alpha^*/2} < 1.5263$$

• So for any alpha such that  $z_{\alpha/2} < 1.5263$ , we will be  $100 * (1 - \alpha)$  percent confident that the true change was greater than zero. How do we find this  $\alpha$ ? Well, if

$$z_{rac{lpha}{2}} = -\Phi^{-1}\left(rac{lpha}{2}
ight)$$
, then  $\Phi\left(-z_{rac{lpha}{2}}
ight) = rac{lpha}{2}$ , and  $lpha = 2\Phi\left(-z_{rac{lpha}{2}}
ight)$ .

- So in this particular case,  $\alpha = 2\Phi(-1.5263)$ .
  - To find this alpha, we now type pnorm(-1.5263) in R, which is the CDF of the standard
     Normal evaluated at its argument. This returns .063.
  - Thus  $\alpha/2 = .063$  and alpha is thus .126.
  - And thus if we are working with confidence intervals of 100 \* (1 .126) = 87.4% or smaller, we will conclude that there was true movement between the two polls.

### 10.2 Hypothesis Testing

- This way of framing the question motivates a process known as **hypothesis testing**. A hypothesis test consists of four elements:
  - 1. A **null hypothesis about a parameter**, which we write as  $H_0$ .
    - This is typically either what the "conventional wisdom" says is the the value of the parameter—or that the parameter is equal to zero.
  - 2. An alternative hypothesis about the parameter,  $H_A$ .
    - This is typically that the parameter is equal to *something different* than the null hypothesis. It may be more specific: that the parameter is either greater than or less than the null hypothesis.
  - 3. A test statistic derived from an estimator of the parameter.
  - 4. A rejection region.
    - The RR specifies the range of values of the test statistic for which the null  $H_0$  is to be *rejected* in favor of the alternative  $H_A$ .
- Choosing the rejection region:
  - RR's are associated with two kinds of error:
    - \* Type I error (a.k.a. a "false positive") is made if  $H_0$  is rejected when  $H_0$  is actually true.
      - ·  $Pr(\text{Type I error}) = \alpha$ . (Yes, the very same  $\alpha$  we've been working with.)
    - \* Type II error (a.k.a. a "false negative") is made if  $H_0$  is accepted when  $H_A$  is actually true.
      - ·  $Pr(\text{Type II error}) = \beta$ .
  - $\alpha$  and  $\beta$  are two very practical ways to measure the goodness of a statistical test. We call  $\alpha$  the test's **level of significance**. We call the quantity  $1 \beta$  the test's **statistical power**. In the best of all worlds, we want a test's level of significance to be low and its power to be high. In reality, we always face a tradeoff between these two goals.

 To illustrate this tradeoff, consider the data from which we constructed the earlier CI about Trump and Biden. Let's re-pose this question in terms of a hypothesis test, where

$$H_0: p_1 - p_2 = 0$$

$$H_A : p_1 - p_2 \neq 0$$

- Here our *test statistic* is the difference between our two sample proportions,  $\hat{p}_1 \hat{p}_2$ . And our rejection region includes the values of the statistic for which we reject the null for our chosen  $\alpha$ .
  - \* Here, the rejection region are those values of  $\hat{p}_1 \hat{p}_2$  for which the constructed CI does not include zero. This would lead us to say (with  $100 * (1 \alpha)$ % confidence) that the change between the two polls was greater than zero.
  - \* What is this region? Let's look at our CI again:

$$(\widehat{p}_1 - \widehat{p}_2) \pm z_{\alpha/2} \sqrt{\frac{\widehat{p}_1(1-\widehat{p}_1)}{n_1} + \frac{\widehat{p}_2(1-\widehat{p}_2)}{n_2}}.$$

\* Had  $(\widehat{p}_1 - \widehat{p}_2)$  been big enough that  $(\widehat{p}_1 - \widehat{p}_2) - z_{\alpha/2} \sqrt{\frac{\widehat{p}_1(1-\widehat{p}_1)}{n_1} + \frac{\widehat{p}_2(1-\widehat{p}_2)}{n_2}} > 0$ , then our CI would not have incorporated zero. That is, if

$$\begin{array}{ccc} (\widehat{p}_{1}-\widehat{p}_{2}) & > & z_{\alpha/2}\sqrt{\frac{\widehat{p}_{1}(1-\widehat{p}_{1})}{n_{1}}+\frac{\widehat{p}_{2}(1-\widehat{p}_{2})}{n_{2}}} \\ \\ \frac{(\widehat{p}_{1}-\widehat{p}_{2})}{\sqrt{\frac{\widehat{p}_{1}(1-\widehat{p}_{1})}{n_{1}}+\frac{\widehat{p}_{2}(1-\widehat{p}_{2})}{n_{2}}}} & > & z_{\alpha/2}, \end{array}$$

we should reject the null and accept  $H_A$ .

- \* So, a few questions:
- \* what sample sizes would we have needed for our difference in sample proportions  $(\hat{p}_1 \hat{p}_2)$  to have been found statistically different from zero with 95% confidence?

(Assume  $n_1 = n_2 = n$ .)

$$\frac{3.1}{\sqrt{\frac{(55.6)(100-55.6)}{n} + \frac{(52.5)(100-52.5)}{n}}} > 1.96$$

$$\frac{3.1}{\sqrt{\frac{4962.4}{n}}} > 1.96$$

$$\frac{3.1}{1.96} > \sqrt{\frac{4962.4}{n}}$$

$$\left(\frac{3.1}{1.96}\right)^2 > \frac{4962.4}{n}$$

$$n > 1,983.7$$

\* We would have needed two samples of at least 1,984 in size.

## 10.3 Hypothesis tests vis-a-vis confidence intervals

- Now let's pose the following question:
  - The conventional wisdom is that our parameter  $\theta$  is equal to a certain value. Specifically the null hypothesis says  $\theta = \theta_0$ .
  - I have theoretical reasons to believe that  $\theta \neq \theta_0$ . This is my alternative hypothesis.
  - I've obtained a point estimate  $\hat{\theta}$  that is not equal to  $\theta_0$ .
  - This, of course, does not in itself destroy the null hypothesis. Why? Because it's
    possible that the null is true and I happened to get this very different estimate from the
    null simply by chance.
  - This leads to a key question:
- Having obtained a point estimate  $\hat{\theta} \neq \theta_0$ , how sure am I now that  $\theta$  is not equal to  $\theta_0$ ?
  - This is the question we ask when we conduct *hypothesis tests*.
  - Often this value is  $\theta_0 = 0$ , but in practice it can be any value.
  - Regardless of the value of  $\theta_0$ , we can fully define the distribution of our estimator,  $\hat{\theta}$  if  $\theta_0$  is true.

- We know that CLT tells us that the standardized version of *any* estimator  $\widehat{\theta}$  that is a linear combination of i.i.d. random variables is distributed Normal in large samples:  $\frac{\widehat{\theta} \mu_{\widehat{\theta}}}{\sigma_{\widehat{\theta}}} N(0,1)$ .
- of course  $\mu_{\widehat{\theta}} = E\left(\widehat{\theta}\right)$  and if  $\widehat{\theta}$  unbiased and if  $\theta_0$  is true then by definition  $E\left(\widehat{\theta}\right) = \theta_0$ .
- So now rewrite as  $\frac{\widehat{\theta}-\theta_0}{\sigma_{\widehat{\theta}}}$   $\sim N\left(0,1\right)$
- And this is the distribution of the standardized version of our estimator  $\hat{\theta}$  "under the null." It is centered around 0, but due to *chance variation* in the sampling process individual estimates are closer or further away from 0 in a pattern described by the standard Normal density.
- [Draw another density curve.] Sometimes it's easier to envision this as the unstandardized version of  $\widehat{\theta}$ . It is centered about  $\theta_0$ , but due to *chance variation* in the sampling process individual estimates are closer or further away from  $\theta_0$  in a pattern described by the Normal density with variance  $\sigma_{\widehat{\theta}}^2$ .

# 10.4 A Two-Tailed Hypothesis Test

- With these tools in place, we can conduct hypothesis tests.
- Recall that such tests consist of:
  - 1. A **null hypothesis about a parameter**, which we write as  $H_0$ .
    - This is typically either what the "conventional wisdom" says is the the value of the parameter—or that the parameter is equal to zero.
  - 2. An alternative hypothesis about the parameter,  $H_A$ .
    - This is typically that the parameter is equal to *something different* than the null hypothesis. It may be more specific: that the parameter is either greater than or less than the null hypothesis.
  - 3. A test statistic derived from an estimator of the parameter.
  - 4. A rejection region.
    - The RR specifies the range of values of the test statistic for which the null  $H_0$  is to be *rejected* in favor of the alternative  $H_A$ .

- We begin by picking a level of confidence,  $\alpha$ . Recall that this is the probability of a Type I error, that is Pr (reject  $H_0|H_0$  true). Typically in our discipline this number is .05, or 5 percent.
- We then look at the standard Normal density curve, and identify the range of extreme values of  $\widehat{\theta}$  that we will observe  $\alpha$  percent of the time in repeated sampling.
- These extreme values are those greater than  $z_{\frac{\alpha}{2}}$  and those less than  $-z_{\frac{\alpha}{2}}$ , where  $\Phi\left(-z_{\frac{\alpha}{2}}\right)=\frac{\alpha}{2}$ .
- Values in these ranges are the **rejection region** for our test. If  $\widehat{\theta}$  falls in the rejection region, we reject  $H_0: \theta = \theta_0$  in favor of  $H_A: \theta \neq \theta_0$ .
- If  $\widehat{\theta}$  does not fall in the rejection region, we fail to reject  $H_0: \theta = \theta_0$ .
- In practice, we are usually working with the unstandardized version of our estimator, and so:
  - We reject  $H_0$  if  $\widehat{\theta} < \theta_0 z_{\frac{\alpha}{2}}\sigma_{\widehat{\theta}}$  or if  $\widehat{\theta} > \theta_0 + z_{\frac{\alpha}{2}}\sigma_{\widehat{\theta}}$ .
  - Otherwise, we fail to reject  $H_0$ .

# 10.5 A One-Tailed Hypothesis Test

- Now consider a case where we have a stronger alternative hypothesis.
- Specifically, our hypothesis is **signed**. Rather than  $H_A: \theta \neq \theta_0$ , I have theoretical reason to claim say,  $H_A: \theta > \theta_0$
- How does this change things?
- We again pick a level of confidence,  $\alpha$ , again typically 5 percent.
- We then look at the standard Normal density curve, and identify the range of values of  $\widehat{\theta}$  greater than  $\theta_0$  that we will observe  $\alpha$  percent of the time in repeated sampling.
- Draw ONE-TAILED HYPOTHESIS TEST diagram on board

# 10.6 Cooking the books

- So here's the thing. Let's say you have a test statistic (some realization of  $\widehat{\theta}$ ) whose value is greater than  $\theta_0$ . You make a *ex post* ("based on actual results") hypothesis that  $H_A: p_1 p_2 \ge 0$  and conduct a one-tailed hypothesis test. This hypothesis is not based on theory. Are you cooking the books?
  - Yes. Knowing that  $\hat{\theta} > \theta_0$ , to reject  $H_0$  with a one-tailed test, you need

$$\widehat{\theta} > z_{\alpha}$$
.

– But to reject  $H_0$  with a two-tailed test, you need

$$\widehat{\theta} > z_{\frac{\alpha}{2}}.$$

- Typically political scientists are skeptical of one-tailed tests because they can look awfully post hoc. Most of the hypothesis tests you'll see in journals are two-tailed.