# **Biostatistics I: Hypothesis testing**

# **Categorical data: Proportion tests**

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- z-test for proportions
- Bionomial test
- ► Examples

#### Assumptions

- > The observations are independent of one another
- The sample size is large enough to use the normal approximation N(np, np(1 - p))
  - np > 10 and n(1 p) > 10, where n is the number of observations and p the proportion

Is the probability of being diagnosed with asthma now different than it was 50 years ago?

### Hypothesis

 $H_{O}: \pi = \pi_{O}$  $H_{1}: \pi \neq \pi_{O}$ 

# One sample *z*-test for proportions: Theory

# Hypothesis

```
If one-tailed
Is the probability of being diagnosed with asthma now higher
than it was 50 years ago?
H_0: \pi = \pi_0
H_1: \pi > \pi_0
```

#### or

Is the probability of being diagnosed with asthma now lower than it was 50 years ago?  $H_0: \pi = \pi_0$  $H_1: \pi < \pi_0$ 

# One sample *z*-test for proportions: Theory

# Test statistic

For large sample sizes, the distribution of the test statistic is approximately normal

$$Z = \frac{p - \pi_0}{\sqrt{\frac{\pi_0(1 - \pi_0)}{n}}}$$

- Sample proportion: *p*
- Population proportion:  $\pi_0$
- Number of subjects: *n*

If continuity correction is applied:  $z = \frac{p - \pi_0 + c}{\sqrt{\frac{\pi_0(1 - \pi_0)}{n}}}$ ,

where

• 
$$c = -\frac{1}{2n}$$
 if  $p > \pi_0$   
•  $c = \frac{1}{2n}$  if  $p < \pi_0$   
•  $c = 0$  if  $|p - \pi_0| < \frac{1}{2n}$ 

# One sample *z*-test for proportions: Theory

#### **Sampling distribution**

- z-distribution
- Critical values and p-value

### Type I error

• Normally  $\alpha$  = 0.05

#### **Draw conclusions**

Compare test statistic (z) with the critical values<sub>α/2</sub> or the p-value with α

If **one-tailed**: Compare test statistic with the critical value $_{\alpha}$ 

Is the probability of being diagnosed with asthma now different than it was 50 years ago?

### Hypothesis

 $H_{O}: \pi = \pi_{O}$  $H_{1}: \pi \neq \pi_{O}$ 

# Hypothesis

 $H_0: \pi = \pi_0$  $H_1: \pi \neq \pi_0$ 

### **Collect and visualize data**

Х	Freq
No	47
Yes	53

50 years ago we had  $\pi_0$  = 0.6

# Test statistic

(with no continuity correction):

$$Z = \frac{p - \pi_0}{\sqrt{\frac{\pi_0(1 - \pi_0)}{p}}} = \frac{0.53 - 0.6}{\sqrt{\frac{0.6(1 - 0.6)}{100}}} = -1.43$$

**Type I error**  $\alpha = 0.05$ 

### **Critical values**

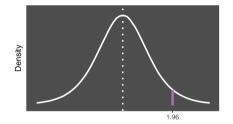
Using R we get the critical values from the z-distribution: critical value<sub> $\alpha/2$ </sub> = critical value<sub>0.05/2</sub>

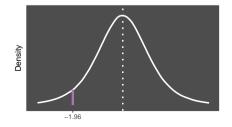
qnorm(p = 0.05/2, lower.tail = FALSE)

```
[1] 1.959964
-critical value<sub>\alpha/2</sub> = -critical value<sub>0.05/2</sub>
```

qnorm(p = 0.05/2, lower.tail = TRUE)

[1] -1.959964





#### **Critical values**

```
If one-tailed
```

```
critical value<sub>\alpha</sub>:
gnorm(p = 0.05, lower.tail = FALSE)
```

```
or
```

```
-critical value<sub>\alpha</sub>:
qnorm(p = 0.05, lower.tail = TRUE)
```

### **Draw conclusions**

We reject the  $H_0$  if:

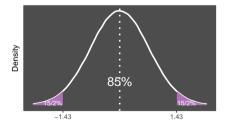
►  $z > \text{critical value}_{\alpha/2}$  or  $z < - \text{critical value}_{\alpha/2}$ 

We have -1.43 > -1.96  $\Rightarrow$  we do not reject the  $H_0$ 

Using  $\mathbb{R}$  we obtain the p-value from the *z*-distribution:

2 \* pnorm(q = -1.43, lower.tail = TRUE)

[1] 0.152717



Is the probability of being diagnosed with asthma in the Netherlands different than in Belgium?

### Hypothesis

 $H_0: \pi_1 = \pi_2$  $H_1: \pi_1 \neq \pi_2$ 

### **Test statistic**

For large sample sizes, the distribution of the test statistic is approximately normal.

Pooled version:  

$$z = \frac{(p_1 - p_2) - 0}{\sqrt{p(1 - p)\left(\frac{1}{n_1} + \frac{1}{n_2}\right)}}$$

Unpooled version:  

$$z = \frac{(p_1 - p_2) - 0}{\sqrt{\frac{p_1(1 - p_1)}{n_1} + \frac{p_2(1 - p_2)}{n_2}}}$$

- Sample proportion of group 1: p1
- ► Sample proportion of group 2: p<sub>2</sub>
- ▶ Number of subjects in group 1: *n*<sub>1</sub>
- Number of subjects in group 2: n<sub>2</sub>
- Total proportion:  $p = \frac{n_1 p_1 + n_2 p_2}{n_1 + n_2}$

### **Test statistic**

If continuity correction is applied:

Pooled version:  

$$z = \frac{(p_1 - p_2) + \frac{F}{2} \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}{\sqrt{p(1 - p) \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}}$$

Unpooled version:  

$$z = \frac{(p_1 - p_2) + \frac{F}{2} \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}{\sqrt{\frac{p_1(1 - p_1)}{n_1} + \frac{p_2(1 - p_2)}{n_2}}}$$

where

# Two sample *z*-test for proportions: Theory

# **Sampling distribution**

- z-distribution
- Critical values and p-value

# Type I error

• Normally  $\alpha$  = 0.05

### **Draw conclusions**

• Compare test statistic (z) with the critical values or the p-value with  $\alpha$ 

Is the probability of being diagnosed with asthma in the Netherlands different than in Belgium?

### Hypothesis

 $H_0: \pi_1 = \pi_2$  $H_1: \pi_1 \neq \pi_2$ 

### **Collect and visualize data**

Table 1: the Netherlands

xl	Freq
No	47
Yes	53

#### **Test statistic**

(with no continuity correction and pooled version):

$$p = \frac{n_1 p_1 + n_2 p_2}{n_1 + n_2} = \frac{100\ 0.53 + 100\ 0.38}{100 + 100} = 0.46$$
$$z = \frac{(p_1 - p_2) - 0}{\sqrt{p(1 - p)\left(\frac{1}{n_1} + \frac{1}{n_2}\right)}} = \frac{0.53 - 0.38}{\sqrt{0.46(1 - 0.46)\left(\frac{1}{100} + \frac{1}{100}\right)}} = 2.13$$

Table 2: Belgium

x2	Freq
No	62
Yes	38

**Type I error**  $\alpha = 0.05$ 

### **Critical values**

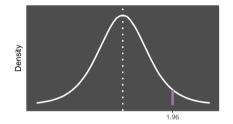
Using R we get the critical values from the z-distribution: critical value<sub> $\alpha/2$ </sub> = critical value<sub>0.05/2</sub>

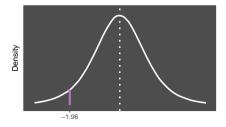
qnorm(p = 0.05/2, lower.tail = FALSE)

```
[1] 1.959964
-critical value<sub>\alpha/2</sub> = -critical value<sub>0.05/2</sub>
```

qnorm(p = 0.05/2, lower.tail = TRUE)

[1] -1.959964





### **Draw conclusions**

We reject the  $H_0$  if:

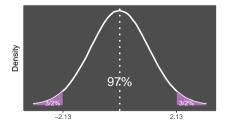
►  $z > \text{critical value}_{\alpha/2}$  or  $z < - \text{critical value}_{\alpha/2}$ 

We have 2.13 > 1.96  $\Rightarrow$  we reject the  $H_0$ 

Using R we obtain the p-value from the *z*-distribution:

2 \* pnorm(q = 2.13, lower.tail = FALSE)

[1] 0.03317161



#### Assumptions

Independent observations

#### Notes..

The binomial test is an exact test

Is the probability of being diagnosed with asthma now different than it was 50 years ago?

### Hypothesis

 $H_{0}: \pi = \pi_{0}$  $H_{1}: \pi \neq \pi_{0}$ 

# **Bionomial test: Theory**

If n is the sample size and k the successes:  $Pr(X = k) = \binom{n}{k}p^k(1-p)^{n-k}$ , where  $\binom{n}{k} = \frac{n!}{k!(n-k)!}$  and ! indicates a factorial

- For any possible outcome of the binomial we obtain the corresponding probability
- We find the p-value by considering the probability of seeing an outcome as, or more, extreme
  - For a one-tailed test,  $H_1: \pi < \pi_0$  $p - value = \Pr(X = 0) + \dots + \Pr(X = k) = \sum_{i=0}^k \Pr(X = i) = \sum_{i=0}^k \binom{n}{i} p^i (1 - p)^{n-i}$
  - Calculating a p-value for a two-tailed test is more complicated, since a binomial distribution is not symmetric if π₀ ≠ 0.5 ⇒ we cannot double the p-value from the one-tailed test

### Type I error

• Normally  $\alpha$  = 0.05

# **Bionomial test: Application**

**Scenario** Is the probability of being diagnosed with asthma now lower than it was 50 years ago?

### Hypothesis

 $H_0: \pi = \pi_0$  $H_1: \pi < \pi_0$ 

# **Collect and visualize data**

- ▶ *n* = 10
- ▶ *k* = 3
- ▶ *p* = 0.3
- $\pi_0 = 0.4$  the probability of being diagnosed with asthma 50 years ago

**P-value** *Pr*(*X* <= 3)

Using R we get the p-value: pbinom(q = 3, size = 10, prob = 0.4)

[1] 0.3822806

#### **Draw conclusions**

We do not reject the  $H_0$ 

# **Bionomial test: Application**

**Scenario** Is the probability of being diagnosed with asthma now higher than it was 50 years ago?

### Hypothesis

 $H_{0}: \pi = \pi_{0}$  $H_{1}: \pi > \pi_{0}$ 

# **Collect and visualize data**

- ▶ *n* = 10
- ▶ *k* = 6
- ▶ *p* = 0.6
- $\pi_0 = 0.4$  the probability of being diagnosed with asthma 50 years ago

### **P-value** Pr(X >= 6) = 1-Pr(X < 6) = 1-Pr(X <= 5)

[1] 0.1662386

#### **Draw conclusions**

We do not reject the  $H_0$