# Module 4. Bivariate analysis: qualitative variables

**Data Science & Al** 

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#### **Learning goals**

- Dependent/independent variable
- Apply suitable analysis techniques for each combination of measurement levels
- Contingency tables and Cramér's V
- Visualization



## Bivariate analysis: overview

Independent	Dependent	Test/Metric
Qualitative	Qualitative	$\chi^2$ -test
		Cramér's <b>V</b>
Qualitative	Quantitative	two-sample <i>t</i> -test
		Cohen's <i>d</i>
Quantitative	Quantitative	_
		Regression, correlation



## **Bivariate analysis**



## **Bivariate analysis**

- ...is determining whether there is an <u>association</u> between two stochastic variables (X and Y).
- Association = you can <u>predict</u> (to some extent) the value of Y from the value of X
  - o X Independent variable
  - o Y Dependent variable
- Important! Finding an association does NOT imply a causal relation!



## **Example research questions**

- Is there a difference in taste preference between two beverage brands?
- Is there a difference in spending at the campus restaurant between students and staff?
- Do smokers die more often of lung cancer than non-smokers?
- Do men and women have a different opinion on a survey question?
- ..

We will use data/rlanders.csv from the Github repo for lab assignments to explore the last question.



## **Contingency tables**



## **Contingency tables**

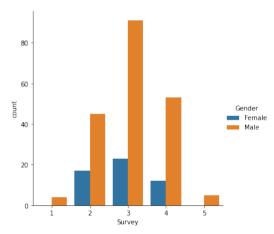
(also: crosstab)

See Python example code in demo-chi-squared

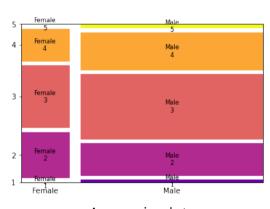
Gender Survey	Female	Male
Strongly disagree	0	4
Disagree	17	45
Neutral	23	91
Agree	12	53
Strongly agree	0	5



#### **Visualization**



A clustered bar chart



A mosaic plot

## **Marginal totals**

Gender Survey	Female	Male	Total
Strongly disagree	0	4	4
Disagree	17	45	62
Neutral	23	91	114
Agree	12	53	65
Strongly agree	0	5	5
Total	52	198	250



#### **Expected values**

If there is no difference (association), we expect the same ratios in each column of the table!

Gender Survey	Female	Male	Total
Strongly disagree	0.832	3.168	4
Disagree	12.896	49.104	62
Neutral	23.712	90.288	114
Agree	13.520	51.480	65
Strongly agree	1.040	3.960	5
Total	52	198	250

In each cell: (row total  $\times$  column total) / n

#### Measuring dispersion

How far is the observed value o from the expected e?

$$\frac{(o-e)^2}{e}$$

Gender Survey	Female	Male
Strongly disagree	0.832	0.219
Disagree	1.306	0.343
Neutral	0.021	0.006
Agree	0.171	0.045
Strongly agree	1.040	0.273



#### The chi-squared statistic

The sum of all these values is notated:

$$\chi^2 = \sum_i \frac{(o_i - e_i)^2}{e_i} \approx 4.255$$

- x is the Greek letter chi
- $o_i$  = number of observations in the i'th cell of the contingency table
- $e_i$  = expected frequency
- Small value ⇒ no association
- Large value ⇒ association



# When is $\chi^2$ large enough?

- $2 \times 2$ -table with  $\chi^2 = 10$ 
  - o Relatively large difference
  - o Indicates association
- $5 \times 5$ -table with  $\chi^2 = 10$ 
  - o Relatively small difference
  - o Does NOT indicate association

We need a metric independent of table size!



#### Cramér's V

$$V = \sqrt{\frac{X^2}{n(k-1)}} = \sqrt{\frac{4.255}{250(2-1)}} \approx 0.130$$

with n the number of observations, k = min(numRows, numCols)

Cramér's V	Interpretation
≈ 0	no association
≈ 0.1	weak association
≈ 0.25	moderate association
≈ 0.5	strong association
≈ 0.75	very strong association
≈ 1	complete association

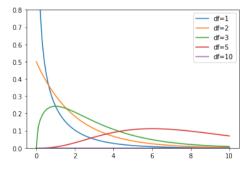


# Chi-squared test for independence



## $\chi^2$ test for independence

- = Alternative to Cramér's V to investigate association between qualitative variables.
- ullet Value of  $\chi^2$  distributed according to the  $\chi^2$  distribution





# $\chi^2$ -distribution in Python

Import scipy.stats For a  $\chi^2$ -distribution with df degrees of freedom:

Function	Purpose
<pre>chi2.pdf(x, df=d) chi2.cdf(x, df=d) chi2.sf(x, df=d) chi2.isf(1-p, df=d)</pre>	Probability density for x  Left-tail probability P(X < x)  Right-tail probability P(X > x)  p% of observations is expected  to be lower than this value  HO  GEN

#### **Test procedure**

- **Step 1.** Formulate hypotheses:
  - o  $H_0$ : there is no association ( $\chi^2$  is "small")
  - o  $H_1$ : there is an association ( $\chi^2$  is "large")
- **Step 2.** Choose significance level, e.g.  $\alpha$  = 0.05
- **Step 3.** Calculate the test statistic,  $\chi^2 = 4.255$



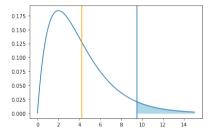
#### **Test procedure (cont.)**

- Step 4. Use  $df = (numRow 1) \times (numCol 1)$  and either:
  - o Determine critical value g so  $P(\chi^2 > g) = \alpha$
  - o Calculate the *p*-value
- Step 5. Draw conclusion:
  - o  $\chi^2 < g$ : do not reject  $H_0$ ;  $\chi^2 > g$ : reject  $H_0$
  - o  $p > \alpha$ : do not reject  $H_0$ ;  $p < \alpha$ : reject  $H_0$



## **Example (Gender vs Survey)**

- g = stats.chi2.isf(0.05, df=4) (result: 9.4877)
- p = stats.chi2.sf(4.2555, df=4) (result: 0.3725)



**Conclusion**: we do not reject the null hypothesis. Differences between expected and observed values are not significantly large. We found no association between variables *Gender* and *Survey* 

## **Test for independence in Python**

SciPy has a function to calculate  $\chi^2$  and p-value from a contingency table:

```
observed = pd.crosstab(rlanders.Survey, rlanders.Gender)
chi2, p, df, expected = stats.chi2_contingency(observed)

print("Chi-squared : %.4f" % chi2)
print("Degrees of freedom: %d" % df)
print("P-value : %.4f" % p)
```



#### **Goodness-of-fit test**



#### Goodness-of-fit test

The  $\chi^2$  test can also be used to determine whether a sample is **representative** for the population.

#### Goodness-of-fit test

This test indicates to what degree a sample corresponds to a null hypothesis regarding the distribution of a qualitative variable over mutually exclusive classes.





Туре	# sample	# population
Mutant	127	35%
Human	75	17%
Alien	98	23%
God	27	8%
Demon	73	17%
Total	400	100%



Is the distribution of the sample (n = 400) representative for the full population (all superheroes)?

- What numbers would you expect if the sample is representative?
- How large are the differences from the *observed* numbers?
  - o small ⇒ distribution is representative
  - o large ⇒ distribution is **not** representative



Is the distribution of the sample (n = 400) representative for the full population (all superheroes)?

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  - o large ⇒ distribution is **not** representative

Can you see the link with contingency tables and Cramer's V?



- Exactly representative ⇒ 35% of superheroes in the sample is a mutant
- The expected number therefore is  $e = 0.35 \times 400 = 140$ .

Therefore:

$$e = n \times \pi$$

If the differences o - e are relatively small they can be attributed to random sampling errors.



Consider  $\chi^2$ :

$$\chi^2 = \sum_{i=1}^n \frac{(o_i - e_i)^2}{e_i}$$

Draw a conclusion based on the value of  $\chi^2$ :

- small ⇒ distribution is representative
- large ⇒ distribution is **not** representative

 $\chi^2$  measures the degree of conflict with the null hypothesis



Superhero type	0	π	е	$\frac{(o-e)^2}{e}$
Mutant	127	35%	140	1.21
Human	75	17%	68	0.72
Alien	98	23%	92	0.39
God	27	8%	32	0.78
Demon	73	17%	68	0.37
			χ <sup>2</sup> =	3.47



- The test statistic  $\chi^2$  follows the  $\chi^2$  distribution.
- Critical value g from the  $\chi^2$  distribution: this is dependent on the number of degrees of freedom (df). In general:

$$df = k - 1$$

with k the number of categories.

• The critical value g for a given significance level  $\alpha$  and number of degrees of freedom df can be calculated in Python using the function isf().

$$P(\chi^2 < g) = 1 - \alpha$$



#### **Testing Procedure**

- Formulate hypotheses
  - o  $H_0$ : sample is representative for the population
  - o  $H_1$ : sample is not representative for the population
- 2. Choose significance level:  $\alpha = 0.05$



#### **Testing Procedure**

1. Calculate test statistic:

$$\chi^2 = \sum_{i=1}^n \frac{(o_i - e_i)^2}{e_i}$$

- 1.1 **Critical area**: Calculate g so that  $P(\chi^2 < g) = 1 \alpha$
- 1.2 **Probability value**: Calculate  $p = 1 P(X < \chi^2)$
- 2. Conclusion (the test is always right-tailed):
  - 2.1  $\chi^2 < g \Rightarrow$  do not reject  $H_0$ ,  $\chi^2 > g \Rightarrow$  reject  $H_0$
  - 2.2  $p > \alpha \Rightarrow$  do not reject  $H_0$ ,  $p < \alpha \Rightarrow$  reject  $H_0$



- g = stats.chi2.isf(0.05, df=4) (result: 9.4877)
- p = stats.chi2.sf(3.4679, df=4) (result: 0.4828)

**Conclusion.**  $\chi^2 \approx 3.47 < g \approx 9.4877$ , so we don't reject the null hypothesis. This sample is representative for the population.



## **Goodness-of-fit test in Python**

```
observed = np.array([127, 75, 98, 27, 73])
   expected p = np.array([.35, .17, .23, .08, .17])
   expected = expected p * sum(observed)
3
   chi2, p = stats.chisquare(f obs=observed, f exp=expected)
   print("\chi^2 = %.4f" \% chi2)
   print("p = %.4f" % p)
```



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#### Standardized residuals



#### **Example: families**

Consider all families with exactly 5 children in a given community.

#### **Example: families**

Consider all families with exactly 5 children in a given community. When we look at the number of boys/girls, there are 6 possible combinations:

- 1. 5 boys
- 2. 4 boys, 1 girl
- 3. 3 boys, 2 girls
- 4. 2 boys, 3 girls
- 5. 1 boy, 4 girls
- 6. 5 girls

A survey was conducted regarding 1022 families with exactly 5 children

Are the observed numbers in the 6 classes representative for a population in which the probability of having a boy is equal to the probability of having a girl, or more concrete 0.5?

i	0	1	2	3	4	5
o <sub>i</sub>	58	149	305	303	162	45



i	0	1	2	3	4	5	_
o <sub>i</sub>	58	149	305	303	162	45	

If the assumption is true, the probability  $\pi_i$  to have i boys is determined by a binomial distribution with parameters n=5 and p=0.5. For example, the probability to have 2 boys out of 5 children is equal to:

$$(0.5)^2 \times (1 - 0.5)^{5-2} \times {5 \choose 2}$$

In general (you don't have to know why):

$$\pi_i = {5 \choose i} \times 0.5^i \times 0.5^{5-i} = \frac{5!}{i!(5-i)!} \times 0.5^i$$



i	0	1	2	3	4	5	Total
o <sub>i</sub>	58	149	305	303	162	45	1022
$\pi_i$	0,031	0,156	0,313	0,313	0,156	0,031	1
$e_i$	31,9	159,7	319,4	319,4	159,7	31,9	1022
<u>(o−e)²</u> e	21,268	0,715	0,647	0,840	0,033	5,343	28,846
r <sub>i</sub> •	4,686	-0,921	-0,970	-1,105	0,199	2,348	



- Formulate both hypotheses
  - o  $H_0$ : the sample is representative for the population
  - o  $H_1$ : the sample is not representative for the population
- 2. **Determine**  $\alpha$  **and** n:  $\alpha$  = 0.01 and n = 1022.
- 3. Value of the test statistic in the sample:

$$\chi^2 = \sum_{i=1}^n \frac{(o_i - e_i)^2}{e_i} \approx 28.846$$

4. Calculate and plot critical area: The critical value is 15.0863. Our test statistic is inside the critical area, so we can reject  $H_0$ .

#### **Standardized Residuals**

#### Standardized Residuals

The standardized residuals indicate which classes make the greatest contribution to the value of  $\chi^2$ .

$$r_i = \frac{o_i - n\pi_i}{\sqrt{n\pi_i(1 - \pi_i)}}$$

- $r_i \in [-2, 2] \Rightarrow$  "acceptable" values
- $r_i < -2 \Rightarrow$  underrepresented
- $r_i > 2 \Rightarrow$  overrepresented

**Conclusion:** families in which all children are of the same gender are overrepresented.

#### Cochran's rules



#### **Conditions**

In order to apply the  $\chi^2$ -test, the following conditions must be met (Rule of Cochran)

- 1. For all categories, the expected frequency e must be greater than 1.
- 2. In a maximum of 20 % of the categories, the expected frequency  $\underline{e}$  may be less than 5.

