# Chi-Square test calculator

### Goodness of fit test, Test of independence, McNemar test

<u>Video</u> <u>Information</u> <u>Chi-squared test for variance</u> <u>Chi-Square Calculator</u>

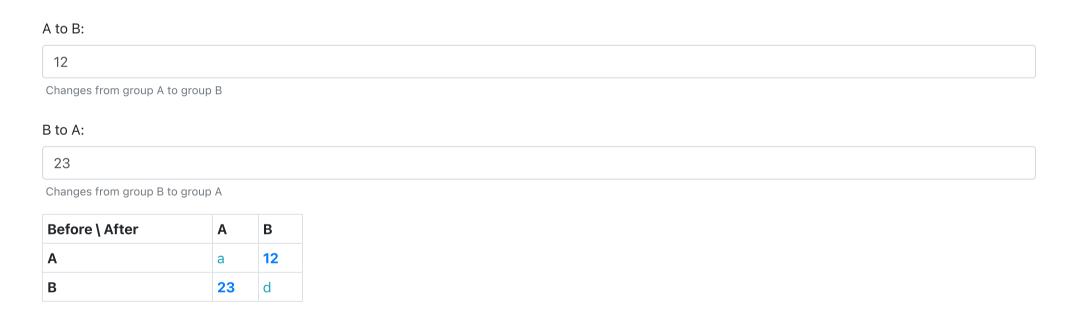
### Test calculation

**Right-tailed** - for the goodness of fit test, the test of independence / the test for association, or the McNemar test, you can use only the right tail test. The calculator includes results from the Fisher calculator, binomial test, McNemar Mid-p, simulation.

right (H₁: after ≥ before)				<b>~</b>
Test:	Goodness of fit	Name:	Population	
Significance level (α):	0.05	m:	0	
Effect:	Medium	Effect size (w):	0.3	
Continuity correction:	False	Simulation repeats:	100000	
Digits:	4			
The continuity correction tends to  Step by step	overcorrect and increase the possibility of a <u>type II error</u> . Hence	ee the resulting p-value is usually <b>too</b>	large. It's controversial, so we recommend to leave it as fals	е
<ul><li>Enter raw data directly</li><li>Enter raw data from exce</li></ul>	I			

### McNemar chi-square test

Matched pairs



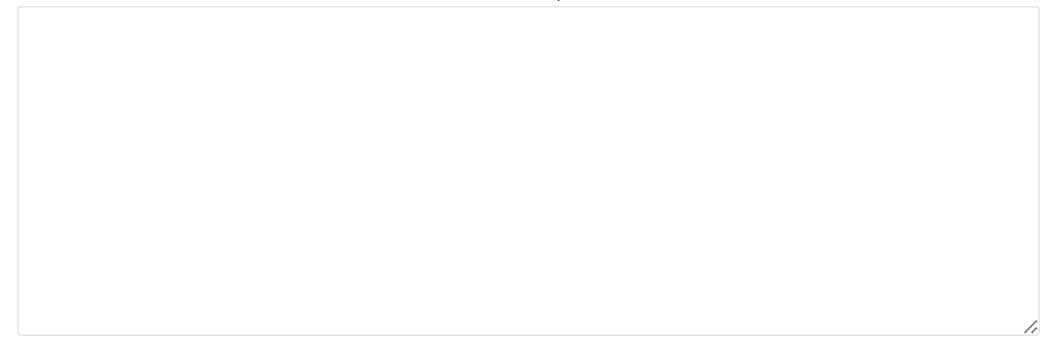
### Enter sample data

Categories //	Conventional //	Assertiveness-based //
Low	3 2 2 1 1 2	311111
Medium	2 2 3 2 2 3	2 2 2 2 3 2
High	455355	455545

### Enter sample data

You may copy data from Excel, Google sheets or any tool that separate data with **Tab** and **Line Feed**.

Copy the data, one block of 3 consecutive columns includes the top header row and left header column, and paste below. example



It is okay to leave empty cells, empty cells or non numeric cells won't be counted

#### \*Choose Expected Frequencies or Expected Probabilities.

The sum of the **expected values** must be equal to the sum of the observed frequencies or to one.



How to do with R?

### Results

19/07/2022, 14:33

The expected values should be at least 1 for all the values, and at least 5 for 80% of the values Blue - less than 5, but still okay.

Red - potential issue.

$$\chi^{2} = \frac{(NaN-NaN)^{2}}{NaN} + \frac{(NaN-NaN)^{2}}{NaN} + \frac{(NaN-NaN)^{2}}{NaN} = NaN$$

P-value = 1 - p(  $\chi^2(2) \le NaN$  ).









#### 

#### <u>. H<sub>0</sub> hypothesis</u>

Since p-value  $< \alpha$ ,  $H_0$  is rejected.

he statistical model does not fit the observations

### <u>!. P-value</u>

The p-value equals NaN, ( $p(x \le \chi^2) = NaN$ ). It means that the chance of type I error (rejecting a correct H<sub>0</sub>) is small: NaN (NaN%). The smaller the p-value the more it supports H<sub>1</sub>.

#### 3. The statistics

The test statistic  $\chi^2$  equals NaN, which is not in the 95% region of acceptance: [- $\infty$  : 5.9915].

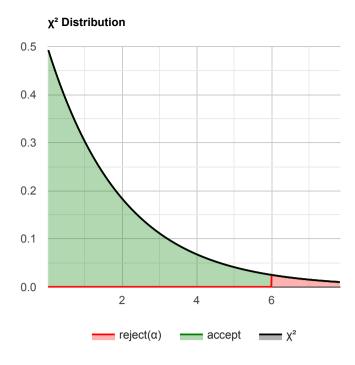
#### I. Effect size

The observed effect size phi is large, NaN. This indicates that the magnitude of the difference between the observed data and the expected data is large.

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### More tests

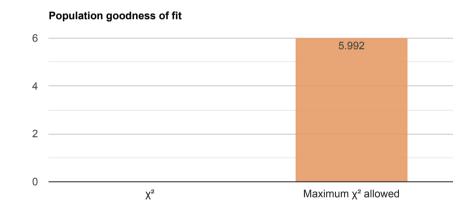
#### What test should I choose?

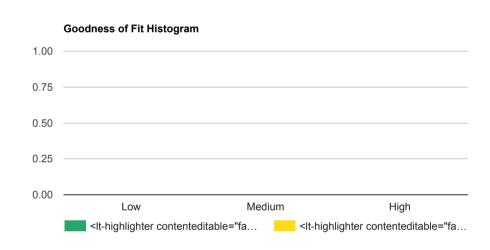
#### **Simulation test**

The simulation run 100,000 times, the duration is 0.1 seconds. Running more times will have better accuracy, but the default should be sufficient. Since the simulation is random, each run will have a slightly different result.

P-value equals 0. It means that the chance of type1 error (rejecting a correct H<sub>0</sub>) is small: 0 (0%)

The smaller the p-value, the more it supports  $H_1$ .





### **Test validation**

The requested test was calculated, it is likely you chose the right test.

• <u>Test power</u> +

Although the priori power is low (NaN), the H<sub>0</sub> is rejected.

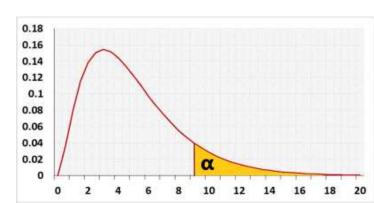
## Information

Hypotheses

Test statistic

χ² distribution

H<sub>0</sub>: Model Fits H<sub>1</sub>: Model Doesn't Fit



Target: Check if the statistical model fits the observations.

The test uses Chi-square distribution.

### McNemar chi-square test

The test checks only the cases when the status of the dichotomous variable was changed.

The null assumption is that the probability to switch from A to B equals the probability to switch from B to A, equals 0.5.

Before \ After	Α	В
Α	No change	A to B
В	B to A	No change

The number of cases with no change, A to A or B to B, doesn't change the result of the McNemar test.

### Chi-square test for independence

The null assumption is that the two categorical variables are independent.

### R Code

The following R code should produce the same results:

obs<-c(3 2 2 1 1 2,2 2 3 2 2 3,4 5 5 3 5 5) prob<-c(3 1 1 1 1 1,2 2 2 2 3 2,4 5 5 5 4 5) prob = prob / sum(prob) chisq.test(x=obs, y=NULL, correct = FALSE, p=prob)

#Compute p-values by Monte Carlo simulation chisq.test(x=obs, y=NULL, correct=FALSE, p=prob, simulate.p.value = TRUE, B = 100000)

Goodness of fit example: checking a fair dice.

Model: the probability of each side is equal - 1/6.

H<sub>0</sub>: fair dice. H<sub>1</sub>: unfair dice.

The **groups** are the dice's numbers (1,2,3,4,5,6).

In this example, you throw the dice n times.

**Expected frequencies** - for each group are n/6.

**Observed frequencies** - the actual times each number appears.