# Manual for web application p-uniform\*

https://rvanaert.shinyapps.io/p-uniformstar
Author: Robbie C.M. van Aert, R.C.M.vanAert@tilburguniversity.edu

Meta-analysis based on *p*-uniform\* can be conducted by means of this web application.<sup>1</sup> *P*-uniform\* is a method for estimating effect size and between-study variance, testing the null hypothesis of no effect and no between-study variance, and testing for publication bias while taking into account publication bias. The *p*-uniform\* method is based on the statistical theory that the distribution of *p*-values is uniform conditional on the population effect size. See van Aert and van Assen (2018) for more information.

The web application can currently handle three different effect size measures: one-sample mean, two-independent means, and raw correlation coefficients. The method can also be applied to effect sizes that are computed by the user (see below for more information).

### Step-by-step guide for using the web application:

- 1) Select the effect size measure in the meta-analysis (one-sample mean, two-independent means, or one raw correlation) or select User-specified effect sizes.
- 2) Select the alpha level which is used in the primary studies. The default alpha level is .05. Note that *p*-uniform\* assumes that two-tailed tests were conducted in the primary studies. If one-tailed hypothesis tests were used in the primary studies, the alpha level has to be multiplied by two before applying *p*-uniform\*.
- 3) Select whether the effect sizes in the primary studies are in the right-tail of the distribution (i.e., positive effects) or in the left-tail of the distribution (i.e., negative effects).
- 4) Three different methods are currently implemented for the p-uniform\* method. The ML method refers to maximum likelihood estimation of the effect size and the between-study variance. Profile likelihood confidence intervals are computed by means of inverting the likelihood-ratio test. Likelihood-ratio tests are used for testing the null hypotheses of no effect and no between-study variance. The ML method is the recommended method for applying p-uniform\*.

The two other methods (P and LNP) are moment based estimators. The method P is based on the distribution of the sum of independent uniformly distributed random variables (Irwin-Hall distribution) and the LNP method refers to Fisher's method (1950, Chapter 4). The test of no effect is not available for the methods P and LNP.

<sup>&</sup>lt;sup>1</sup> This web application is based on the R package puniform which can be downloaded from GitHub by running the following lines of code in R: devtools::install\_github("RobbievanAert/puniform"); library(puniform)

5) Upload the data as a comma separated file (CSV) with a comma as separator and a period indicating the decimal point. The first row should *exactly* match the names below that depend on the effect size measure that has been selected:

### For <u>one-sample mean</u> based on *t*-statistics:

- tobs → t-statistic of each primary study
- ni → sample size of each primary study

## For <u>one-sample mean</u> based on descriptive statistics:

- mi → group mean of each primary study
- $ni \rightarrow$  sample mean of each primary study
- sdi → standard deviation of each primary study

### For <u>two-independent means</u> based on *t*-statistics:

- tobs  $\rightarrow$  t-statistic of each primary study
- n1i → sample size in group 1 of each primary study
- $n2i \rightarrow$  sample size in group 2 of each primary study

### For two-independent means based on descriptive statistics:

- *m1i* → mean in group 1 of each primary study
- $m2i \rightarrow$  mean in group 2 of each primary study
- $n1i \rightarrow$  sample size in group 1 of each primary study
- n2i → sample size in group 2 of each primary study
- $sd1i \rightarrow$  standard deviation in group 1 of each primary study
- $sd2i \rightarrow$  standard deviation in group 2 of each primary study

#### For one correlation:

- $-ri \rightarrow$  raw correlation coefficient of each primary study
- ni → sample size of each primary study

#### For user-specified effect sizes:

- $-yi \rightarrow$  effect size of each primary study
- vi → sampling variance of each primary study

For example, the CSV file for two-independent means should like like:

#### For *t*-statistics:

	Α	В	С	
1	tobs	n1i	n2i	
2	4	200	200	
3	4	200	200	
-				

# For descriptive statistics:

			~	U	E	F
1 m	<b>1</b> i	m2i	n1i	n2i	sd1i	sd2i
2	0.2	0	200	200	1	1
3	0.2	0	200	200	1	1

6) By clicking the "Analyze" button *p*-uniform is applied to your data.

### References

van Aert, R. C. M., & van Assen, M. A. L. M. (2023). Correcting for publication bias in a meta-analysis with the *p*-uniform\* method. Manuscript submitted for publication.