

# Supplementary material for Quantifying replicability and consistency in systematic reviews

The package `metarep` is an extension to the package `meta`, which allows incorporating replicability-analysis tools to quantify consistency and replicability of treatment effect estimates in a meta-analysis. The tool was proposed by Jaljuli et. al. (submitted) for the fixed-effect and for the random-effects meta-analyses, with or without the common-effect assumption.

## Packages Instalation:

Currently, `metarep` can be downloaded from GitHub since is being processed by CRAN, therefore make sure that the package `devtools` is installed. `metarep` also requires the latest version of `meta` ( $\geq 4.11-0$ , available on [github](#) )

Run the following commands in *console* to install the packages:

```
devtools::install_github( "guido-s/meta"      , force=T )
devtools::install_github( "IJaljuli/metarep", force=T )
```

## Examples:

Here we demonstrate the approach implemented with `metarep` on several examples from the Systematic Review Cochrane Library. These examples are detailed in the paper as well, along with a demonstration of a way to incorporate our suggestions in standard meta-analysis reporting system.

We begin with an example based on fixed-effects meta-analysis from review number CD002943: the effect of mammogram invitation on attendance during the following 12 months.

### 1<sup>st</sup> Example: Review CD002943

```
library(metarep)
data(CD002943_CMP001)

m2943 <- meta::metabin( event.e = N_EVENTS1, n.e = N_TOTAL1,
                       event.c = N_EVENTS2, n.c = N_TOTAL2,
                       studlab = STUDY, comb.fixed = T , comb.random = F,
                       method = 'Peto', sm = CD002943_CMP001$SM[1],
                       data = CD002943_CMP001)
```

```
m2943
```

	OR	95%-CI	%W(fixed)
Sutton-1994	1.2836	[0.9933; 1.6589]	32.3
Somkin-1997	1.8739	[1.5372; 2.2844]	54.2
Turnbull-1991	3.5709	[1.9225; 6.6326]	5.5
Mohler-1995	1.8764	[0.5272; 6.6788]	1.3
Bodiya-1999	1.0758	[0.6110; 1.8941]	6.6

```

Number of studies combined: k = 5

              OR              95%-CI      z  p-value
Fixed effect model 1.6564 [1.4317; 1.9164] 6.78 < 0.0001

Quantifying heterogeneity:
tau^2 = 0.0882 [0.0000; 1.5129]; tau = 0.2970 [0.0000; 1.2300];
I^2 = 70.3% [24.4%; 88.3%]; H = 1.84 [1.15; 2.93]

Test of heterogeneity:
      Q d.f. p-value
13.47   4  0.0092

Details on meta-analytical method:
- Peto method
- DerSimonian-Laird estimator for tau^2
- Jackson method for confidence interval of tau^2 and tau

```

summary(m2943)

```

Number of studies combined: k = 5

              OR              95%-CI      z  p-value
Fixed effect model 1.6564 [1.4317; 1.9164] 6.78 < 0.0001

Quantifying heterogeneity:
tau^2 = 0.0882 [0.0000; 1.5129]; tau = 0.2970 [0.0000; 1.2300];
I^2 = 70.3% [24.4%; 88.3%]; H = 1.84 [1.15; 2.93]

Test of heterogeneity:
      Q d.f. p-value
13.47   4  0.0092

Details on meta-analytical method:
- Peto method
- DerSimonian-Laird estimator for tau^2
- Jackson method for confidence interval of tau^2 and tau

```

The replicability-analysis results follow.

```

m2943.ra <- metarep(x = m2943 , u = 2 , common.effect = F ,t = 0.05 ,report.u.max = T)
m2943.ra

```

	OR	95%-CI	%W(fixed)
Sutton-1994	1.2836	[0.9933; 1.6589]	32.3
Somkin-1997	1.8739	[1.5372; 2.2844]	54.2
Turnbull-1991	3.5709	[1.9225; 6.6326]	5.5
Mohler-1995	1.8764	[0.5272; 6.6788]	1.3
Bodiya-1999	1.0758	[0.6110; 1.8941]	6.6

```

Number of studies combined: k = 5

```

```

              OR              95%-CI      z  p-value
Fixed effect model 1.6564 [1.4317; 1.9164] 6.78 < 0.0001

```

```

Quantifying heterogeneity:

```

```
tau^2 = 0.0882 [0.0000; 1.5129]; tau = 0.2970 [0.0000; 1.2300];
I^2 = 70.3% [24.4%; 88.3%]; H = 1.84 [1.15; 2.93]
```

Test of heterogeneity:

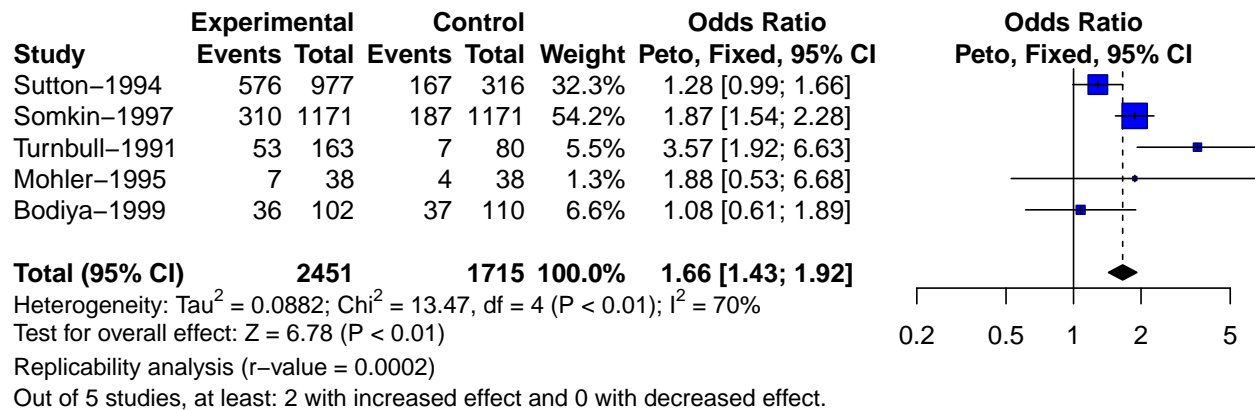
```
Q d.f. p-value
13.47 4 0.0092
```

Details on meta-analytical method:

- Peto method
- DerSimonian-Laird estimator for  $\tau^2$
- Jackson method for confidence interval of  $\tau^2$  and  $\tau$
- replicability analysis (r-value =  $2e-04$ )
- out of 5 studies, at least: 2 with increased effect and 0 with decreased effect.

metarep allows adding replicability results to the conventional forest plots by meta. This can be done by simply applying `meta::forest()` on a `metarep` object.

```
metarep::forest(m2943.ra, layout='revman5', digits.pval = 2, test.overall = T)
```



## 2<sup>nd</sup> Example: Review CD007077

The second example is based on a fixed-effects meta analysis in review CD007077. The main objective of this review is to determine whether PBI/APBI is equivalent to or better than conventional or hypo-fractionated whole breast radiotherapy (WBRT) after breast-conservation therapy for early-stage breast cancer. The primary outcome was Cosmesis.

```
data(CD007077_CMP001)
```

```
m7077 <- meta::metabin( event.e = N_EVENTS1, n.e = N_TOTAL1,
  event.c = N_EVENTS2, n.c = N_TOTAL2,
  studlab = STUDY, comb.fixed = T, comb.random = F,
  method = 'MH', sm = CD007077_CMP001$SM[1],
  data = CD007077_CMP001)
```

```
summary(m7077)
```

Number of studies combined: k = 5

	OR	95%-CI	z	p-value
Fixed effect model	1.5078	[1.1673; 1.9476]	3.14	0.0017

Quantifying heterogeneity:

```
tau^2 = 1.0240; tau = 1.0119; I^2 = 88.4% [75.5%; 94.5%]; H = 2.94 [2.02; 4.26]
```

Test of heterogeneity:

```
Q d.f. p-value
34.47 4 < 0.0001
```

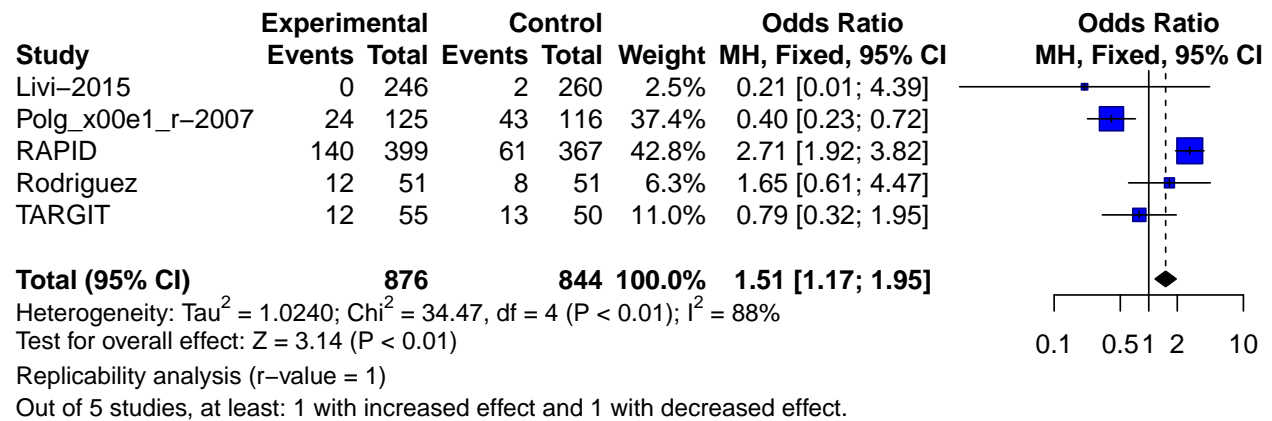
Details on meta-analytical method:

- Mantel-Haenszel method
- DerSimonian-Laird estimator for  $\tau^2$
- Mantel-Haenszel estimator used in calculation of Q and  $\tau^2$  (like RevMan 5)
- Continuity correction of 0.5 in studies with zero cell frequencies

With the replicability analysis:

```
m7077.ra <- metarep(x = m7077 , u = 2, t = 0.05 , report.u.max = T )
```

```
metarep::forest(m7077.ra, layout='revman5',digits.pval = 2 , test.overall = T )
```



### 3<sup>rd</sup> Example:

Based on a random-effects meta-analysis in review CD006823, where the meta-analysis finding was statistically significant. The authors examine the effects of wound drainage after axillary dissection for breast carcinoma on the incidence of post-operative Seroma formation.

```
data(CD006823_CMP001)
```

```
m6823 <- meta::metabin( event.e = N_EVENTS1, n.e = N_TOTAL1,
                        event.c = N_EVENTS2, n.c = N_TOTAL2,
                        studlab = STUDY, comb.fixed = F , comb.random = T,
                        method = 'MH', sm = CD006823_CMP001$SM[1],
                        data = CD006823_CMP001)
```

```
m6823
```

	OR	95%-CI	%W(random)
Cameron-1988	0.1358	[0.0247; 0.7478]	10.0
Somers-1992	0.3341	[0.1633; 0.6837]	20.1
Zavotksy-1998	0.0120	[0.0006; 0.2230]	4.6
Purushotham-2002	0.9457	[0.6305; 1.4183]	23.6
Jain-2004	0.4942	[0.1922; 1.2704]	17.4
Soon-2005	0.6939	[0.0931; 5.1693]	8.1
Classe-2006	1.0446	[0.3665; 2.9774]	16.2

Number of studies combined: k = 7

	OR	95%-CI	z	p-value
Random effects model	0.4576	[0.2293; 0.9130]	-2.22	0.0266

Quantifying heterogeneity:

$\tau^2 = 0.4833$ ;  $\tau = 0.6952$ ;  $I^2 = 67.7\%$  [28.3%; 85.4%];  $H = 1.76$  [1.18; 2.62]

Test of heterogeneity:

Q	d.f.	p-value
18.58	6	0.0049

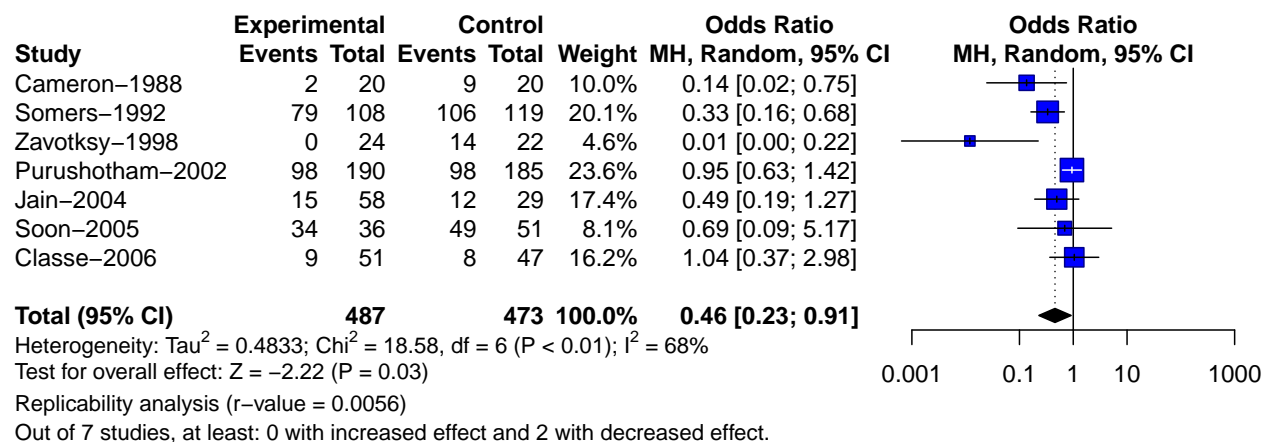
Details on meta-analytical method:

- Mantel-Haenszel method
- DerSimonian-Laird estimator for  $\tau^2$
- Mantel-Haenszel estimator used in calculation of Q and  $\tau^2$  (like RevMan 5)
- Continuity correction of 0.5 in studies with zero cell frequencies

With replicability-analysis:

```
m6823.ra <- metarep(x = m6823 , u = 2, t = 0.05, report.u.max = T )
```

```
metarep::forest(m6823.ra, layout='revman5',digits.pval = 2 , test.overall = T )
```



#### 4<sup>th</sup> Example: Review CD003366

Based on a random-effects meta-analysis in review CD003366. The authors compare chemotherapy regimens on overall effect in Leukopaenia. Pooling 28 studies, the random-effects meta-analysis fails to declare any significant difference between regimens, due to the highly-significant yet contradicting results.

```
data(CD003366_CMP005)
```

```
m3366 <- meta::metabin( event.e = N_EVENTS1, n.e = N_TOTAL1,
                        event.c = N_EVENTS2, n.c = N_TOTAL2,
                        studlab = STUDY, comb.fixed = F , comb.random = T,
                        method = 'MH', sm = CD003366_CMP005$SM[1],
                        data = CD003366_CMP005)
```

m3366

RR	95%-CI	%W(random)
----	--------	------------

ECOG-E1193-x0028_A_x0029_	1.0957	[0.8799; 1.3643]	4.4
EU_x002d_93011	1.4340	[1.1970; 1.7179]	4.8
_x0033_06-Study-Group	1.0824	[1.0196; 1.1490]	5.7
AGO	0.7525	[0.5883; 0.9625]	4.1
Bloher	1.0938	[0.9511; 1.2579]	5.2
Bonneterre	1.3286	[0.8736; 2.0205]	2.6
Bontenbal	1.0568	[0.9505; 1.1749]	5.4
CECOG-BM1	1.1047	[1.0090; 1.2094]	5.6
EORTC-10961	1.0919	[0.9882; 1.2066]	5.5
HERNATA	1.9171	[1.3090; 2.8078]	2.9
Jassem	1.3576	[1.1833; 1.5576]	5.2
Lyman	0.9903	[0.7529; 1.3025]	3.8
Nabholtz	1.1618	[1.0839; 1.2452]	5.7
Rugo	1.4062	[0.3031; 6.5255]	0.3
TRAVIOTA	0.1708	[0.0651; 0.4483]	0.8
_x0033_03-Study-Group	1.0319	[0.9833; 1.0828]	5.8
_x0033_04-Study-Group	0.9988	[0.9499; 1.0501]	5.8
ANZ-TITG	0.4362	[0.3151; 0.6040]	3.3
Dieras	4.1053	[0.9311; 18.1008]	0.4
ECOG-E1193-x0028_B_x0029_	1.2183	[0.9812; 1.5125]	4.4
EORTC-10923	0.4719	[0.3875; 0.5747]	4.6
JCOG9802	1.6019	[1.0899; 2.3545]	2.9
Meier	0.2467	[0.1314; 0.4633]	1.5
Sjostrom	4.8162	[3.2206; 7.2023]	2.7
Talbot	5.7895	[1.4441; 23.2103]	0.4
TOG	0.6048	[0.3018; 1.2119]	1.3
TXT	1.2342	[1.0326; 1.4752]	4.8
Yardley	0.1923	[0.0233; 1.5888]	0.2

Number of studies combined: k = 28

	RR	95%-CI	z	p-value
Random effects model	1.0673	[0.9747; 1.1688]	1.41	0.1596

Quantifying heterogeneity:

tau<sup>2</sup> = 0.0365; tau = 0.1911; I<sup>2</sup> = 89.7% [86.3%; 92.2%]; H = 3.11 [2.70; 3.59]

Test of heterogeneity:

Q	d.f.	p-value
261.41	27	< 0.0001

Details on meta-analytical method:

- Mantel-Haenszel method
- DerSimonian-Laird estimator for tau<sup>2</sup>
- Mantel-Haenszel estimator used in calculation of Q and tau<sup>2</sup> (like RevMan 5)

with replicability-analysis:

```
m3366.ra <- metarep(x = m3366 , u = 2 , t = 0.05, report.u.max = T )
```

```
metarep::forest(m3366.ra, layout='revman5', digits.pval = 2 , test.overall = T )
```

