

The metadat Package: A Collection of Meta-Analysis Datasets for R

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Meta-Analysis Datasets

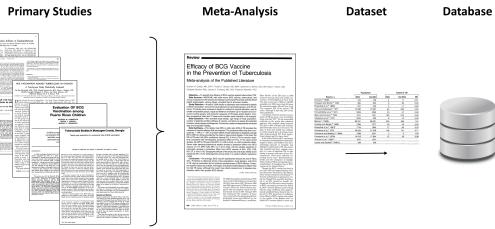
- one of the nice aspects of published meta-analyses: they often (but not always ☺) include the full dataset used
- example: meta-analysis on the effectiveness of the BCG vaccine against tuberculosis (Colditz et al., 1994)

Table 1.—Reports From Clinical Trials Providing Estimates of Efficacy of BCG Vaccine Against Cases of Tuberculosis (TB) and TB Death That Were Used in the Meta-analysis*

Source, Y	Population		Cases of TB		TB Death	
	BCG	No BCG	BCG	No BCG	BCG	No BCG
Aronson ¹¹ 1948	153	309	4	11	0.41	0
Ferguson and Simes ¹⁰ 1949	306	303	6	29	0.20	2
Rosenthal et al. ¹ 1960	231	220	3	11	0.26	0
Hart and Sutherland ¹² 1977	13 598	12 867	62	248	0.24	...
Frimodt-Møller et al. ¹³ 1973	5069	5808	33	47	0.80	...
Stein and Aronson ¹⁴ 1953	1541	1451	160	372	0.65	...
Vandiviere et al. ¹⁵ 1973	2545	629	8	10	0.20	...
Madras ¹⁶ 1989	88 391	88 391	505	499	1.01	...
Coetze and Berjak ¹⁷ 1968	7499	7277	29	45	0.63	...
Rosenthal et al. ¹⁸ 1961 ¹⁸	1716	1665	17	65	0.25	1
Comstock et al. ¹⁹ 1964	50 344	28 558	166	141	0.71	8
Comstock and Webster ²⁰ 1969	2498	2341	5	3	1.56	...
Comstock et al. ²¹ 1976 ²¹	16 913	17 854	27	29	0.98	...
Aronson et al. ¹¹ 1958 ²²	1541	1451	13	68
Levine and Sackett ²³ 1986 ²¹	566	528	8	6
Overall RR (95% confidence interval)						0.49 (0.34-0.70)
						0.29 (0.16-0.53)

Meta-Analysis Datasets

- by extracting the data, one can build up a collection of meta-analysis dataset



- useful for teaching purposes, illustrating/testing meta-analytic methods, validating published analyses, sensitivity checks, ...

Example: BCG Vaccine Meta-Analysis

```

# load the metafor package
library(metafor)

# look at the BCG dataset
dat.bcg

## # trial          author year tpos  tneg  cpos  cneg ablat   alloc
## #  1      Aronson 1948   4  119   11  128   44 random
## #  2 Ferguson & Simes 1949   6  300   29  274   55 random
## #  3 Rosenthal et al 1960   3  228   11  209   42 random
## #  4 Hart & Sutherland 1977  62 13536 248 12619  52 random
## #  5 Frimodt-Møller et al 1973  33 5036  47 5761  13 alternate
## #  6 Stein & Aronson 1953  180 1361  372 1079  44 alternate
## #  7 Vandiviere et al 1973  8 2537  10 619  19 random
## #  8 TPT Madras 1980  505 87886 499 87892  13 random
## #  9 Coetze & Berjak 1968  29 7470  45 7232  27 random
## # 10 Rosenthal et al 1961  17 1699  65 1600  42 systematic
## # 11 Comstock et al 1974  186 50448 141 27197  18 systematic
## # 12 Comstock & Webster 1969  5 2493  3 2338  33 systematic
## # 13 Comstock et al 1976  27 16886 29 17825  33 systematic
  
```

Example: BCG Vaccine Meta-Analysis

```

# calculate log risk ratios and corresponding sampling variances
dat <- escalc(measure="RR", ai=tpos, bi=tneg,
              ci=cpos, di=cneg, data=dat.bcg)
dat

## # trial          author year . yi vi
## #  1      Aronson 1948 . -0.8893 0.3256
## #  2 Ferguson & Simes 1949 . -1.5854 0.1946
## #  3 Rosenthal et al 1960 . -1.3481 0.4154
## #  4 Hart & Sutherland 1977 . -1.4416 0.0200
## #  5 Frimodt-Møller et al 1973 . -0.2175 0.0512
## #  6 Stein & Aronson 1953 . -0.7861 0.0069
## #  7 Vandiviere et al 1973 . -1.6209 0.2230
## #  8 TPT Madras 1980 . 0.0120 0.0040
## #  9 Coetze & Berjak 1968 . -0.4694 0.0564
## # 10 Rosenthal et al 1961 . -1.3713 0.0730
## # 11 Comstock et al 1974 . -0.3394 0.0124
## # 12 Comstock & Webster 1969 . 0.4459 0.5325
## # 13 Comstock et al 1976 . -0.0173 0.0714
  
```

Example: BCG Vaccine Meta-Analysis

```

# random-effects model (using log risk ratios and variances as input)
res <- rma(yi, vi, data=dat, method="DL")

## ## Random-Effects Model (k = 13; tau^2 estimator: DL)
## ## tau^2 (estimated amount of total heterogeneity): 0.3088 (SE = 0.2299)
## ## tau (square root of estimated tau^2 value):      0.5557
## ## I^2 (total heterogeneity / total variability):   92.12%
## ## H^2 (total variability / sampling variability): 12.69
## ##
## ## Test for Heterogeneity:
## ## Q(df = 12) = 152.2330, p-val < .0001
## ##
## ## Model Results:
## ##
## ## estimate      se     zval    pval    ci.lb    ci.ub
## ## -0.7141 0.1787 -3.9952 <.0001 -1.0644 -0.3638
  
```

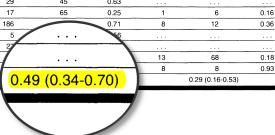
Example: BCG Vaccine Meta-Analysis

```
# predicted average risk ratio (and 95% CI/PI)
predict(res, transf=exp, digits=2)
```

```
## pred ci.lb ci.ub pi.lb pi.us
## 0.49 0.34 0.70 0.16 1.54
```

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Rosenblatt et al. ¹⁶ 1963	233	233	9	11	0.36	0
Hert and Sutherland ¹⁷ 1977	13 598	12 887	62	248	0.24	...
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Stein and Aronson ¹⁹ 1953	1541	1451	186	372	0.46	...
Stein et al. ²⁰ 1974	2453	2459	8	10	0.70	...
Matus et al. ²¹ 1965	88 391	88 391	505	498	1.01	...
Coezee and Bergh ²² 1968	799	7277	29	49	0.63	...
Rosenblatt et al. ¹⁶ 1951	1716	1665	17	65	0.25	1
Comstock et al. ²³ 1974	50 634	27 338	186	186	0.71	8
Comstock et al. ²⁴ 1969	2498	2341	5	10	0.50	...
Comstock et al. ²⁵ 1974	16 913	17 684	27	30	0.89	...
Aronson et al. ²⁶ 1958 ²⁴	1541	1451	13	68
Levine and Sackett ²⁷ 1948 ²¹	566	528	8	8
Overall RR (95% confidence interval)					0.29 (0.16-0.53)	



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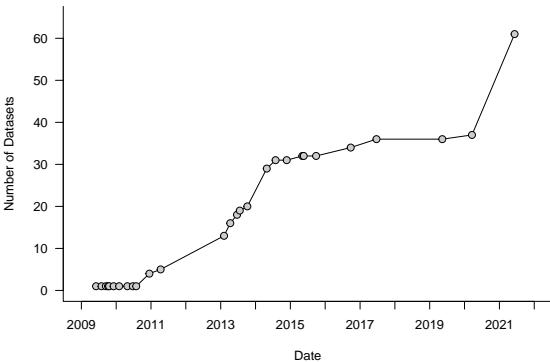
Example: BCG Vaccine Meta-Analysis

```
# sensitivity analyses using different methods/models
[... a whole bunch of code here ...]
print(tab)
```

	pred	ci.lb	ci.ub	tau^2
## Standard Random-Effects Model (DL estimator)	0.49	0.34	0.70	0.313
## Mantel-Haenszel Method (fixed-effects model)	0.64	0.59	0.69	0.000
## Binomial-Normal Model (fixed study effects)	0.49	0.35	0.68	0.271
## Binomial-Normal Model (random study effects)	0.48	0.34	0.68	0.297
## Bayesian Model (bayesmeta package)	0.49	0.32	0.74	0.377
## Bayesian Model (metaBMA package)	0.51	0.36	0.73	0.313
## Bayesian Model (brms package)	0.49	0.32	0.72	0.399

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Number of Datasets Included in the metafor Package



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The metadat Package

- the idea: move datasets into a separate data package
- make it easier to add datasets even without an update to the metafor code and for others to contribute datasets
- started to work on this at the 2019 Evidence Synthesis Hackathon (UNSW Canberra) with Thomas White, Emily Kothe, Daniel Noble, Alistair Senior, and Kyle Hamilton
- first version released on CRAN on August 20th, 2021
- CRAN page: <https://cran.r-project.org/package=metadat>
- Github repo: <https://github.com/wviechtb/metadat>
- documentation: <https://wviechtb.github.io/metadat/>
- currently includes 79 datasets

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Structure

- datasets are called `dat.<author><year>`
- each dataset is documented in a consistent manner
 - a brief/general description of the dataset
 - description of each variable
 - further details about the dataset / meta-analysis
 - the source of the data
 - potentially other relevant references
 - the person who extracted the data
 - examples illustrating use of the data
 - concept terms

Concept Terms

- each dataset is tagged with one or multiple concept terms ...
- in terms of **fields/topics** (e.g., climate change, criminology, ecology, education, oncology, psychology, social work)
- in terms of **outcome measures** (e.g., correlation coefficients, Cronbach's alpha, risk ratios, standardized mean differences)
- in terms of **methods/concepts** (e.g., cluster-robust inference, cumulative meta-analysis, meta-regression, multivariate models, network meta-analysis, outliers, publication bias)
- extremely useful, but only if terms are used consistently
- full list of concept terms used so far: <https://wviechtb.github.io/metadat/reference/datsearch.html#details>

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Search Functionality

- can see all datasets with `help(package=metadat)` or at: <https://wwiechtb.github.io/metadat/reference/index.html>
- `datsearch()` function can be used to search based on concept terms or the full text

```
# find all datasets tagged with 'standardized mean differences'  
datsearch("standardized mean differences")  
  
# find all datasets tagged with 'odds ratio' AND 'multilevel'  
datsearch(c("odds ratio", "multilevel"))  
  
# do a full-text search for the term 'infarct'  
datsearch("infarct", concept=FALSE)
```

Contributing New Datasets

- detailed workflow for contributing new datasets: <https://wwiechtb.github.io/metadat/index.html#contributing-new-datasets>
- some guiding principles:
 - consistent naming scheme for datasets
 - distinguish 'raw data' from the dataset included in the package
 - data preparation script to turn raw data into the `.rda` file
 - `prep_dat()` function to create template for help (`.Rd`) file
- either make pull request via GitHub or just send us the files

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ESMARConf2022 Hackathon

- `datsearch()` is nice but want something more fancy: a Shiny app that replicates this functionality in a more interactive way
- opportunity to add additional datasets to the package, improve the documentation, and consider additional functionality
- results / outcomes to be reported ...

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Thank You for Your Attention!

Questions, Comments, Suggestions?

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🌐 <https://www.wvbauer.com/>

🌐 <https://www.metafor-project.org/>

🐦 [@wwiechtb](#)

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