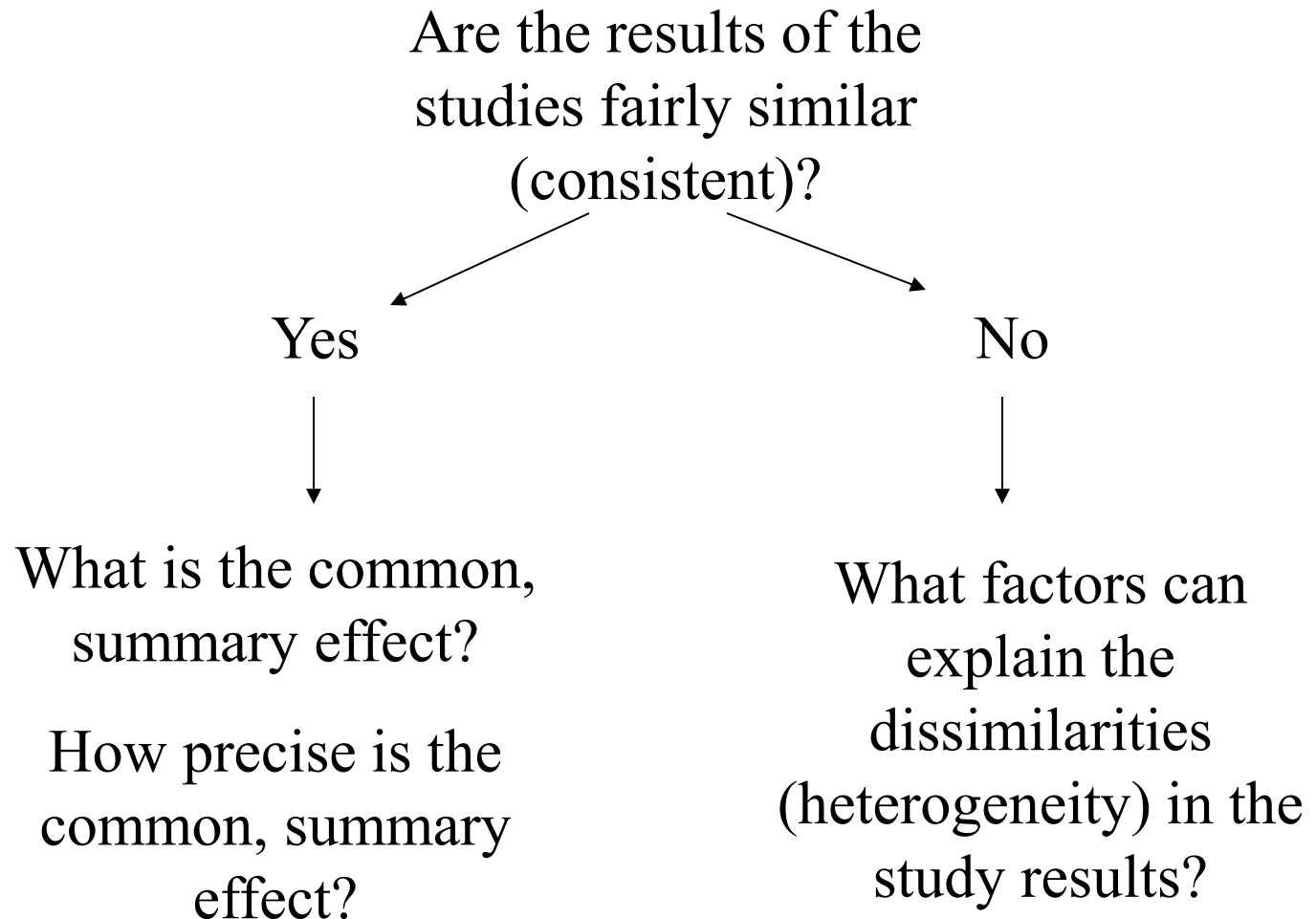


Data Analysis in Systematic Reviews

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Central questions of interest





Steps in data analysis & presentation

1. Tabulate summary data
2. Graph data
3. Check for heterogeneity
4. Perform a meta-analysis if heterogeneity is not a major concern
5. If heterogeneity is found, identify factors that can explain it
6. Evaluate the impact of study quality on results
7. Explore the potential for publication bias

I. Tabulate summary data

- Prepare tables comparing studies with respect to:
 - Year
 - Setting
 - Patients
 - Intervention
 - Comparison
 - Outcome (results)
 - Quality
- Gives a 'first hand' feel for the data
- Can make some assessment of quality and heterogeneity

Tabulate summary data

Example: Cochrane albumin review

Study	Year	Patient population	Intervention	Comparison	Summary measure (RR)	Allocation concealment
Lucas et al.	1978	Trauma	Albumin	No albumin	13.9	Inadequate
Jelenko et al.	1979	Burns	Albumin	Ringer's lactate	0.50	Unclear
Rubin et al.	1997	Hypoalbuminemia	Albumin	No albumin	1.9	Adequate

Cochrane Injuries Group Albumin Reviewers. Human albumin administration in critically ill patients: systematic review of randomised controlled trials. BMJ 1998;317:235-40.

2. Graph summary data

- Efficient way of presenting summary results
- Forest plot:
 - Presents the point estimate and CI of each trial
 - Also presents the overall, summary estimate
 - Allows visual appraisal of heterogeneity
- Other graphs:
 - Cumulative meta-analysis
 - Sensitivity analysis
 - Funnel plot for publication bias
 - Galbraith, L'Abbe plots, etc [rarely used]

Forest Plot

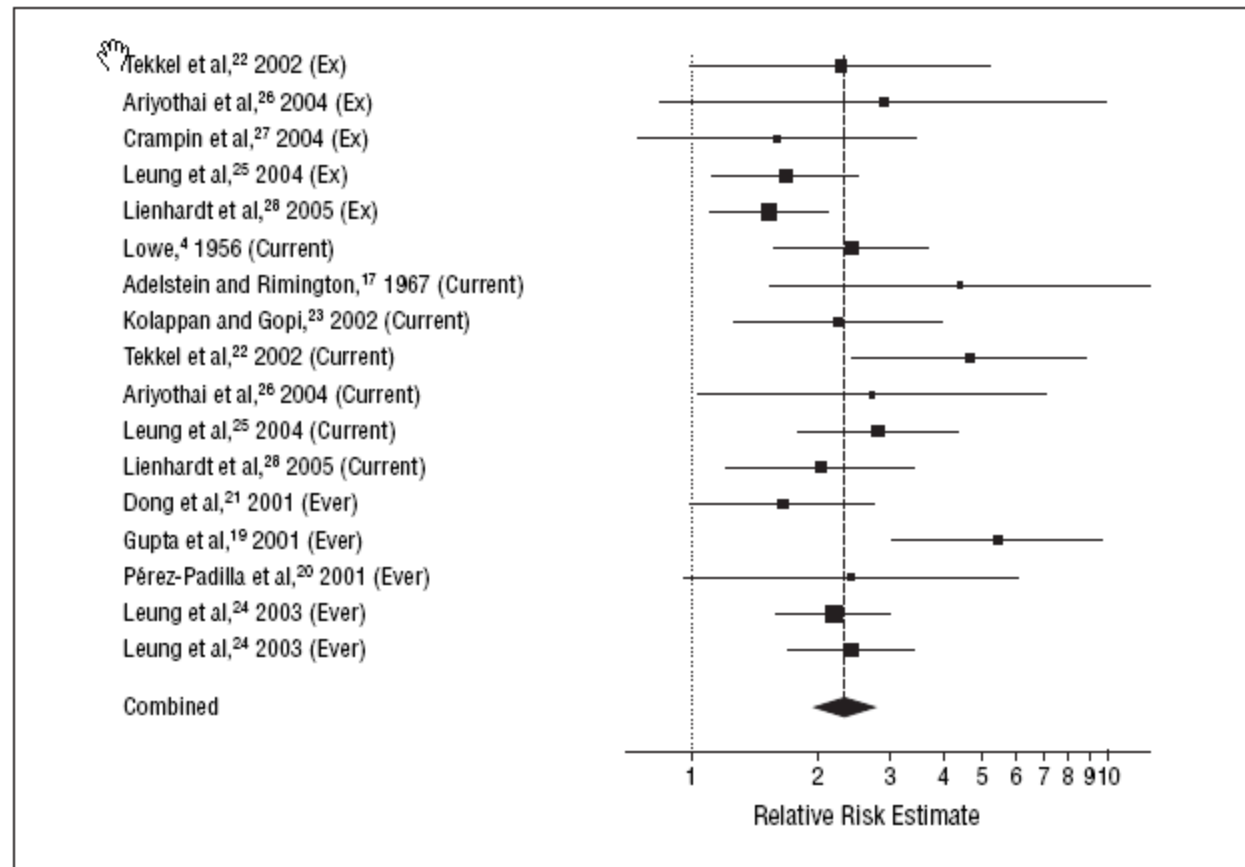


Figure 3. Forest plot of results for men only and for men and women combined in studies^{4,17,19-28} that examined smoking and tuberculosis disease. The smoking type (ex-smokers [Ex], current smokers [Current], and ever smokers [Ever]) of the study population is shown on the y-axis.

Interpreting and understanding meta-analysis graphs

A practical guide

Ideally, clinical decision making ought to be based on the latest evidence available. However, to keep abreast with the continuously increasing number of publications in health research, a primary health care professional would need to read an unsurmountable

meta-analysis before diving into the fine points of the meta-analysis results and drawing conclusions on patient treatment. *Table 1* can guide the assessment.

Meta-analysis graphs



PROFESSIONAL
PRACTICE

Research



Karin Ried

PhD, MSc, GDFH, is Research Fellow & PHCRED Program Manager, Discipline of General Practice, The University of Adelaide, South Australia. karin.ried@adelaide.edu.au

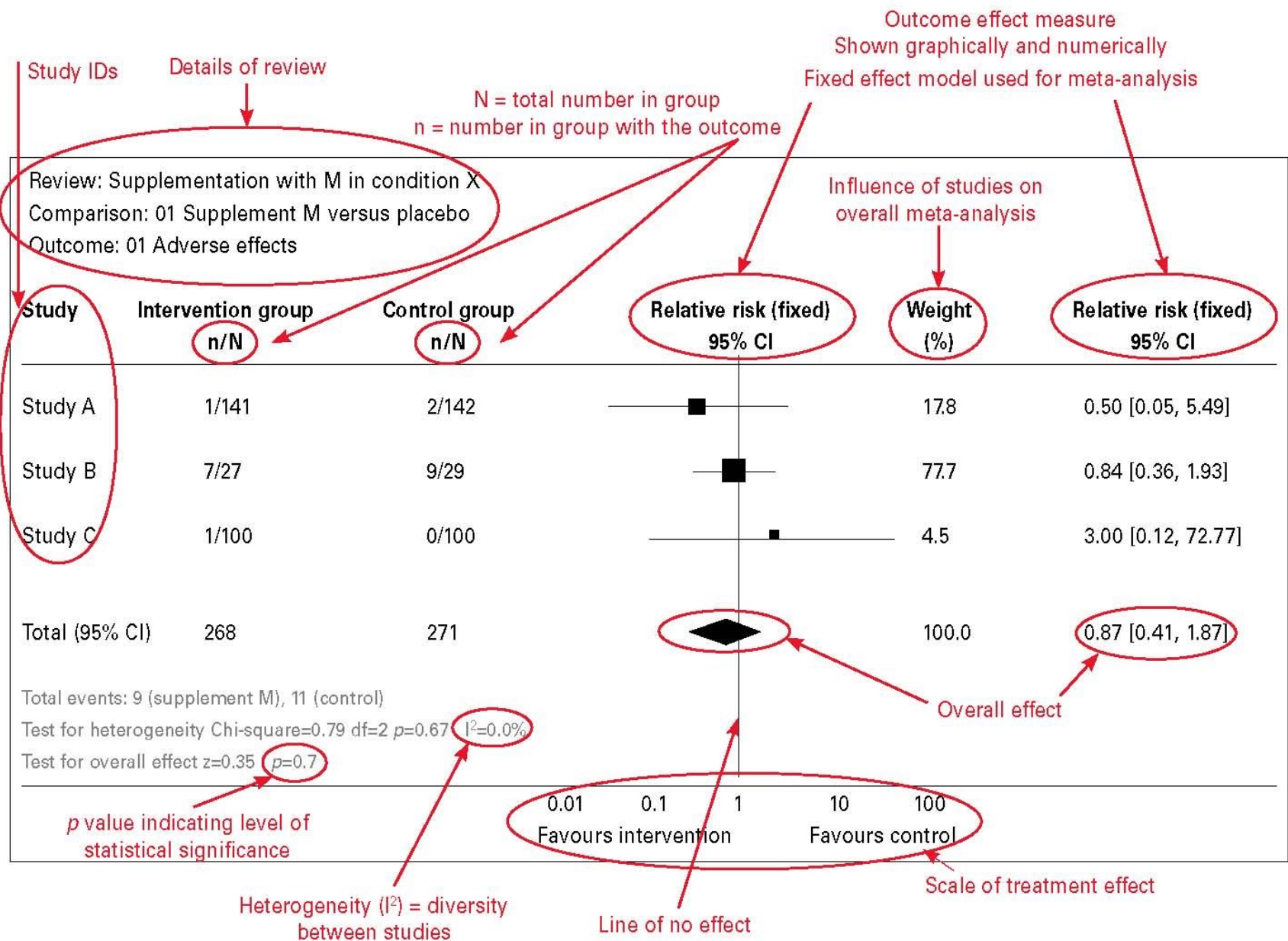


Figure 1. Meta-analysis of binary outcome measure

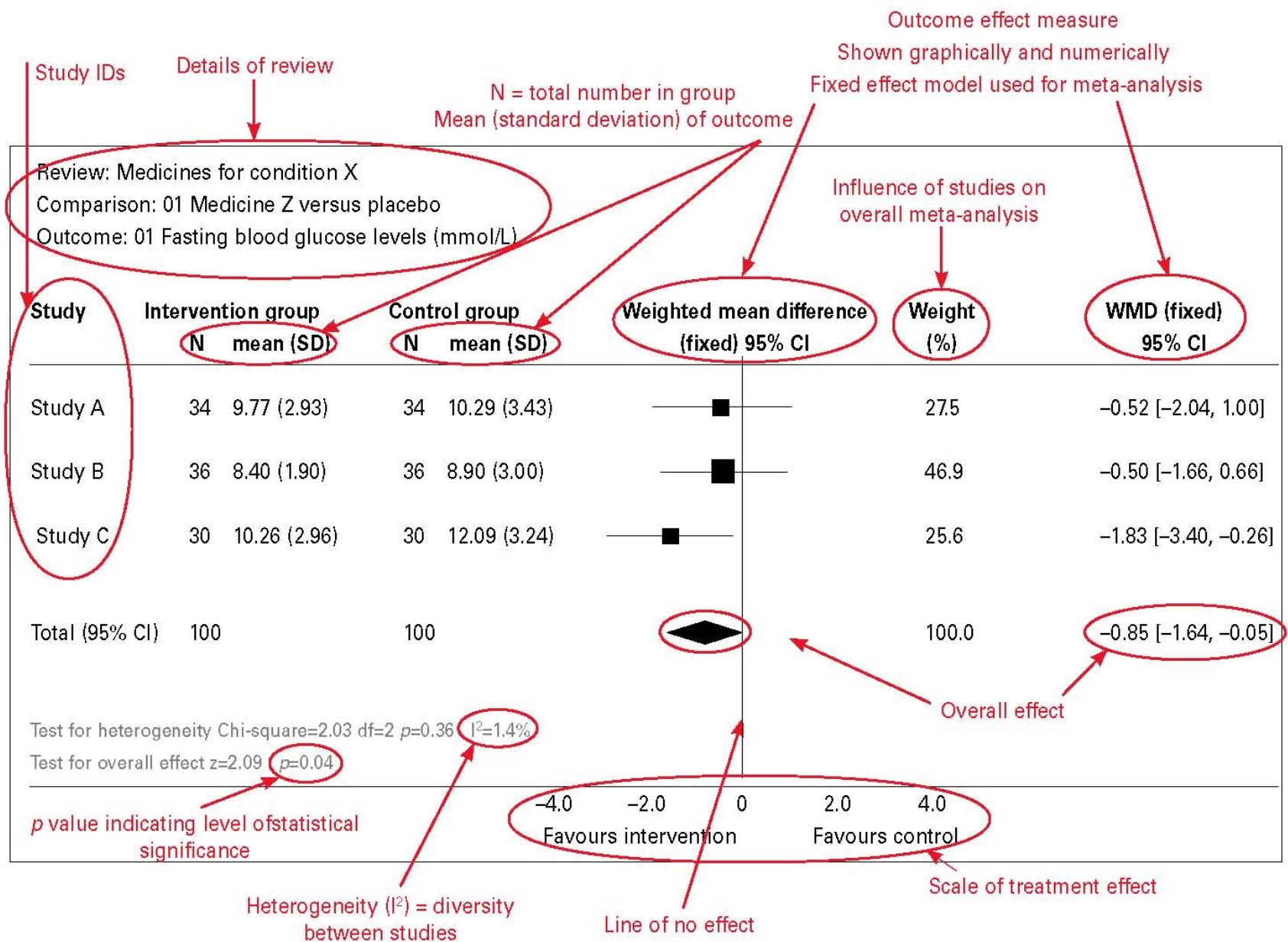
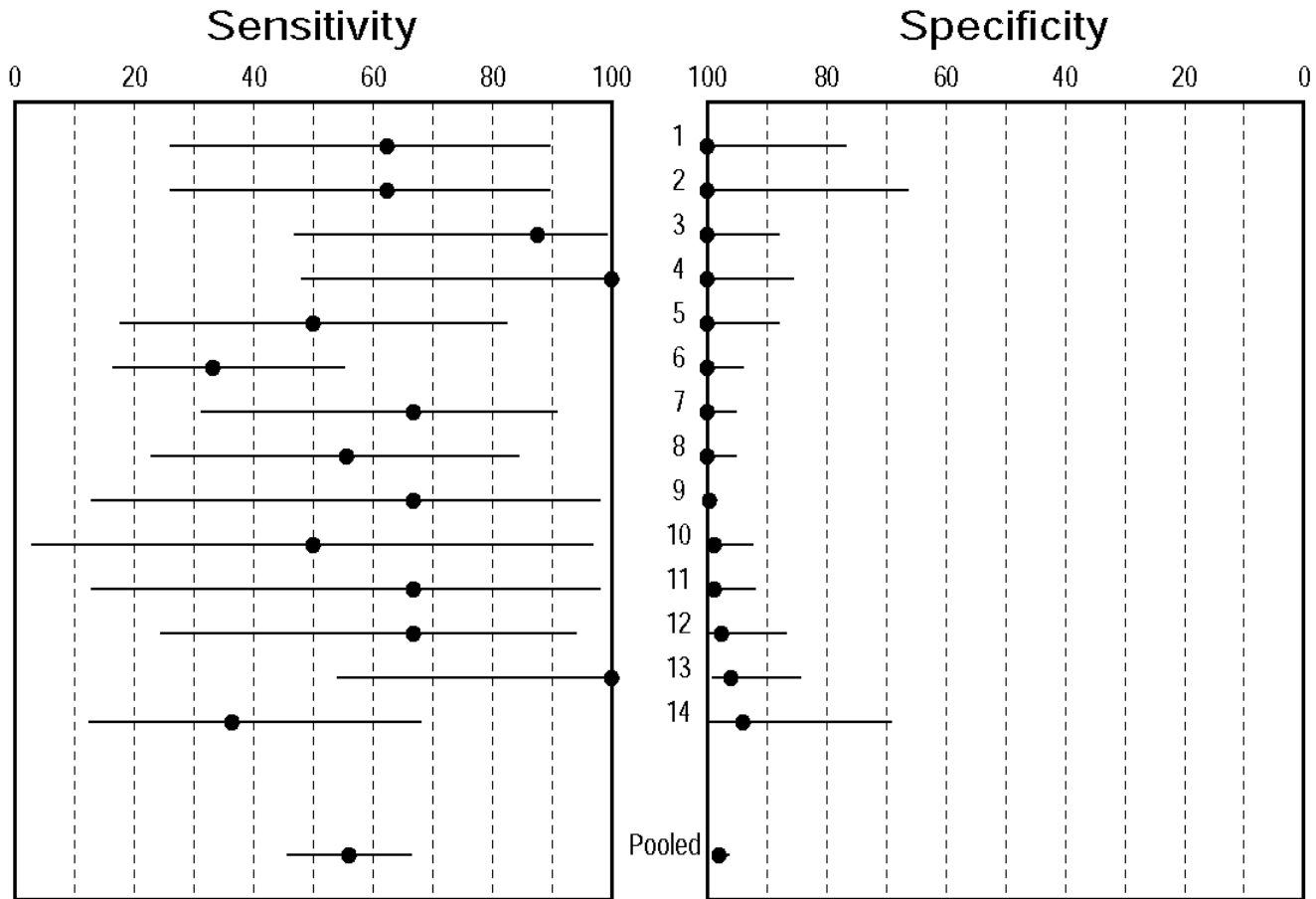


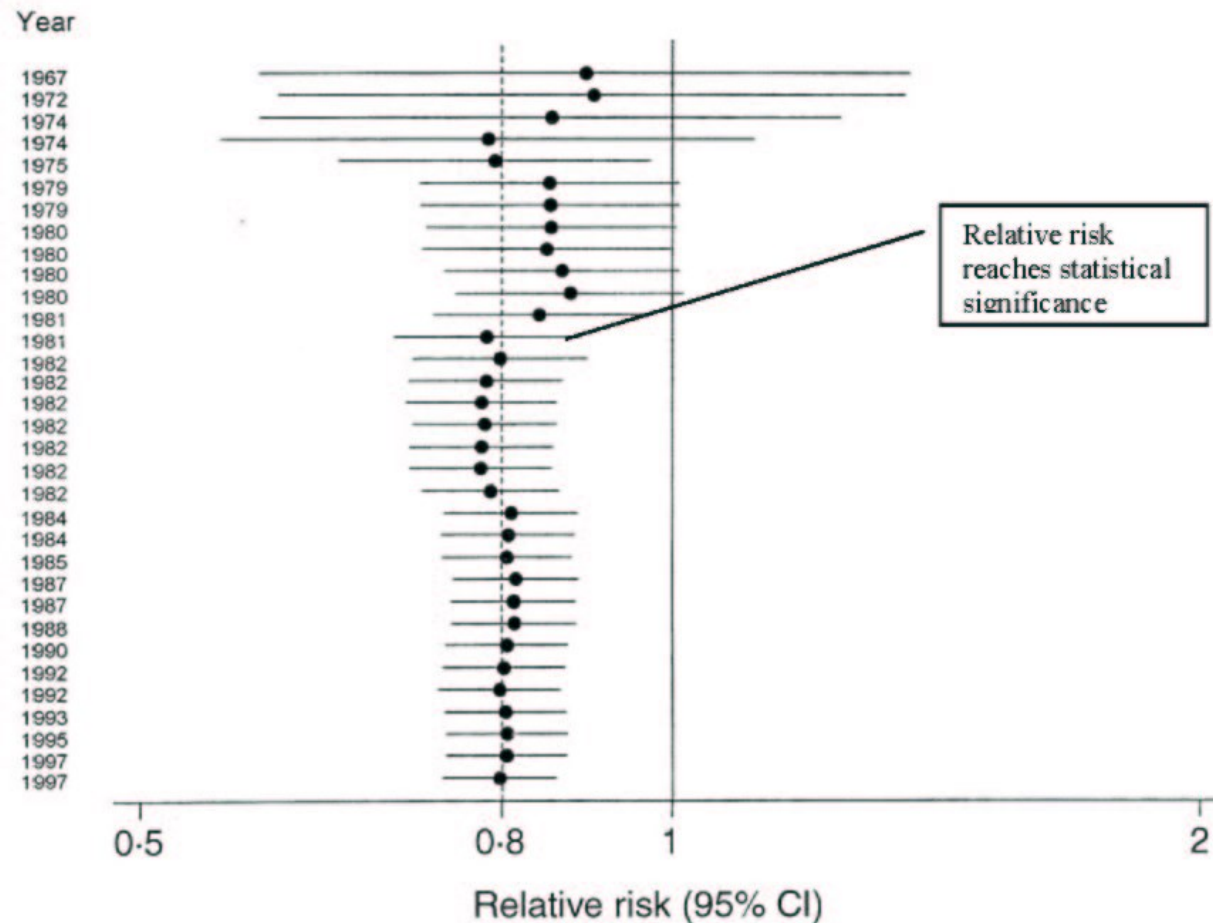
Figure 2. Meta-analysis of continuous outcome measures

Forest Plot: diagnostic studies



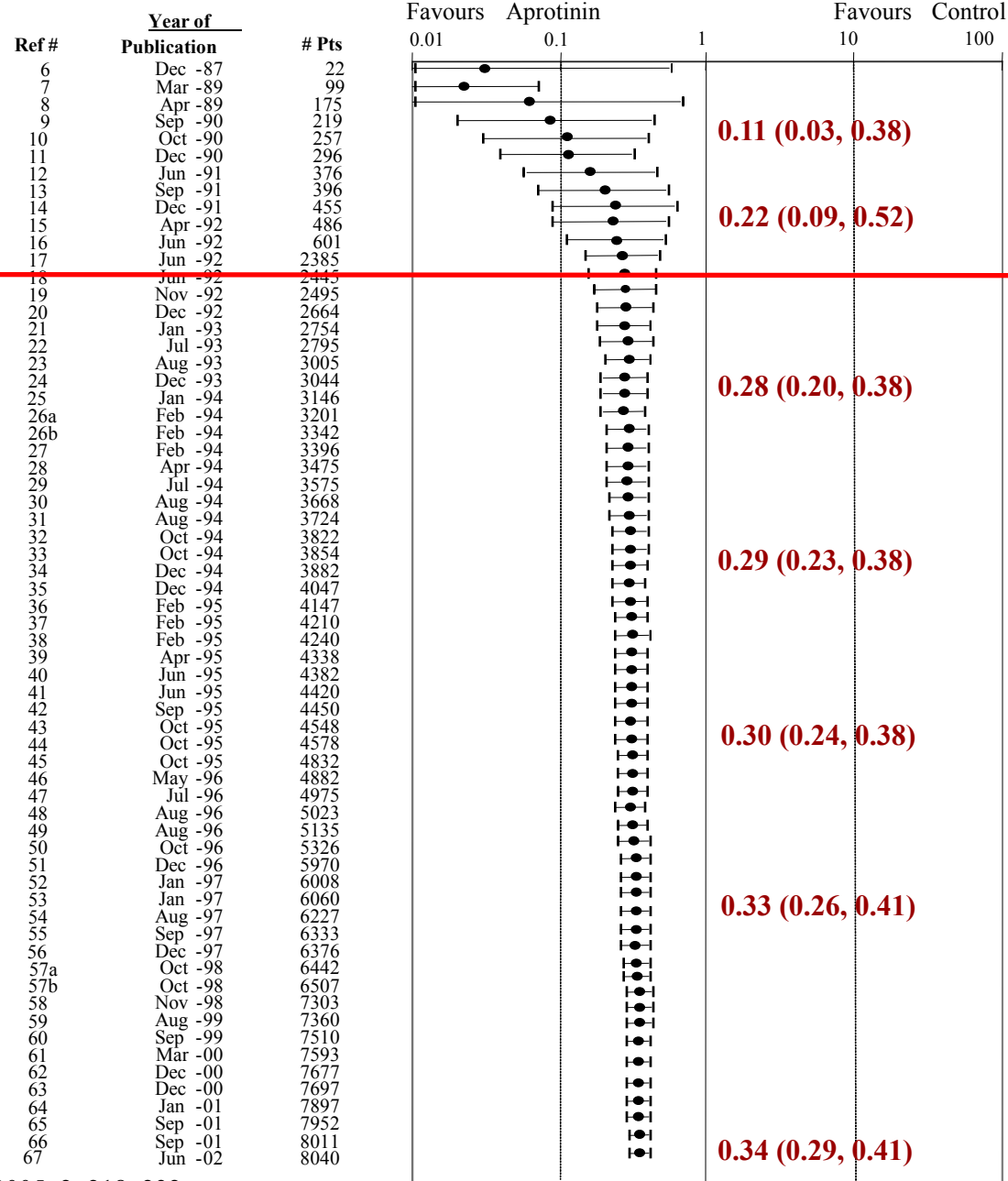
Commercial PCR tests for TB meningitis

Forest Plot: Cumulative Meta-analysis



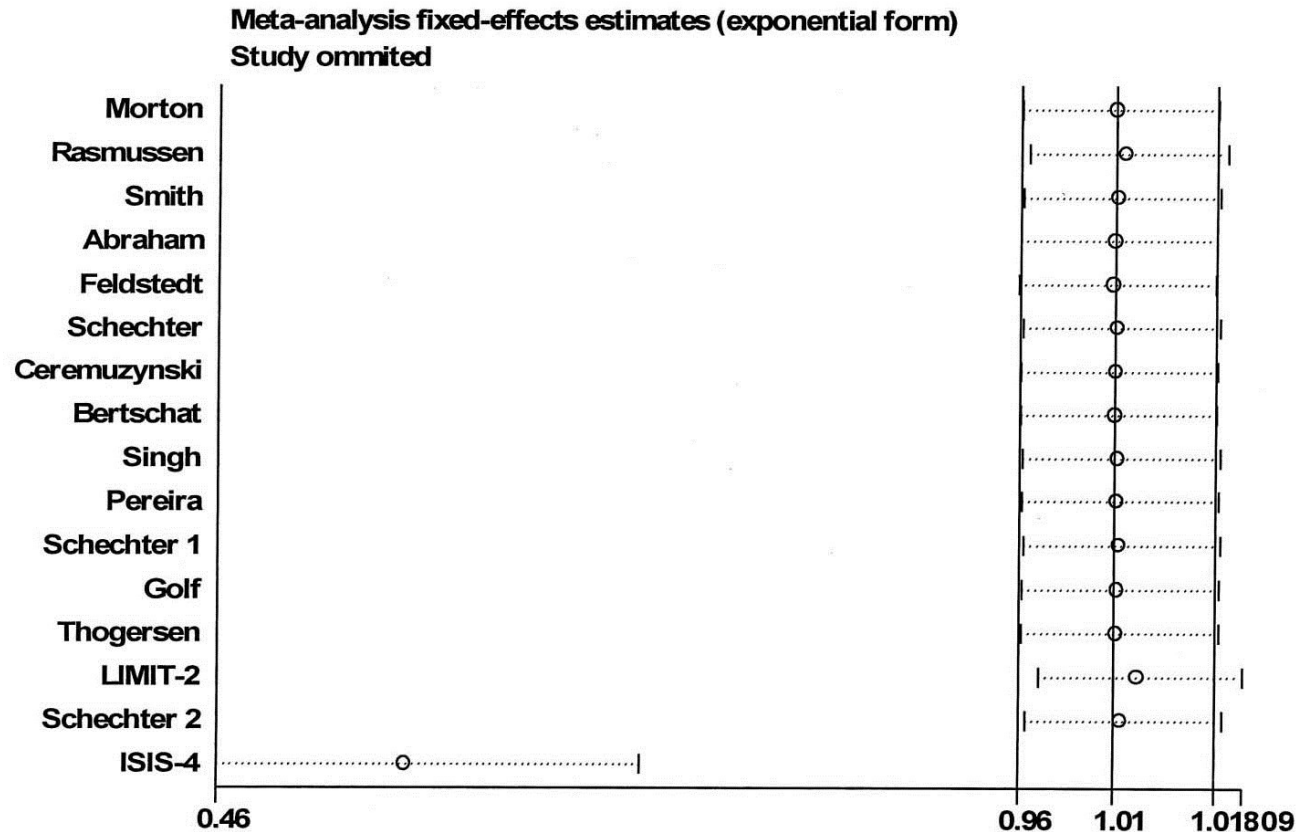
Beta-blockers after acute myocardial infarction

Odds Ratios with 95% Confidence Intervals



Aprotinin
for cardiac
surgery

Sensitivity analysis



IV magnesium for acute myocardial infarction

ISIS-4 trial had >50,000 patients! It showed no survival benefit from the addition of IV magnesium

3. Check for heterogeneity

- Indicates that effect varies a lot across studies
- If heterogeneity is present, a common, summary measure is hard to interpret
- Statistical vs clinical heterogeneity
- Can be due to differences in:
 - Patient populations studied
 - Interventions used
 - Co-interventions
 - Outcomes measured
 - Study design features (eg. length of follow-up)
 - Study quality
 - Random error



Two 'average' men having an 'average' meal.

3. Check for heterogeneity

- How to look for heterogeneity?
 - Visual
 - Forest plot: do confidence intervals of studies overlap with each other and the summary effect?
 - L'Abbe plot
 - Statistical tests:
 - Chi-square test for heterogeneity (Cochran Q test)
 - Tests whether the individual effects are farther away from the common effect, beyond what is expected by chance
 - Has poor power
 - P-value < 0.10 indicates significant heterogeneity
 - I-squared (newly introduced by Higgins et al): % of total variability in effect measure that is attributable to heterogeneity (i.e. not to chance)
 - Values of I-squared equal to 25%, 50%, and 75% representing low, moderate, and high heterogeneity, respectively.

Visual appraisal of heterogeneity

Association between smoking and TB mortality

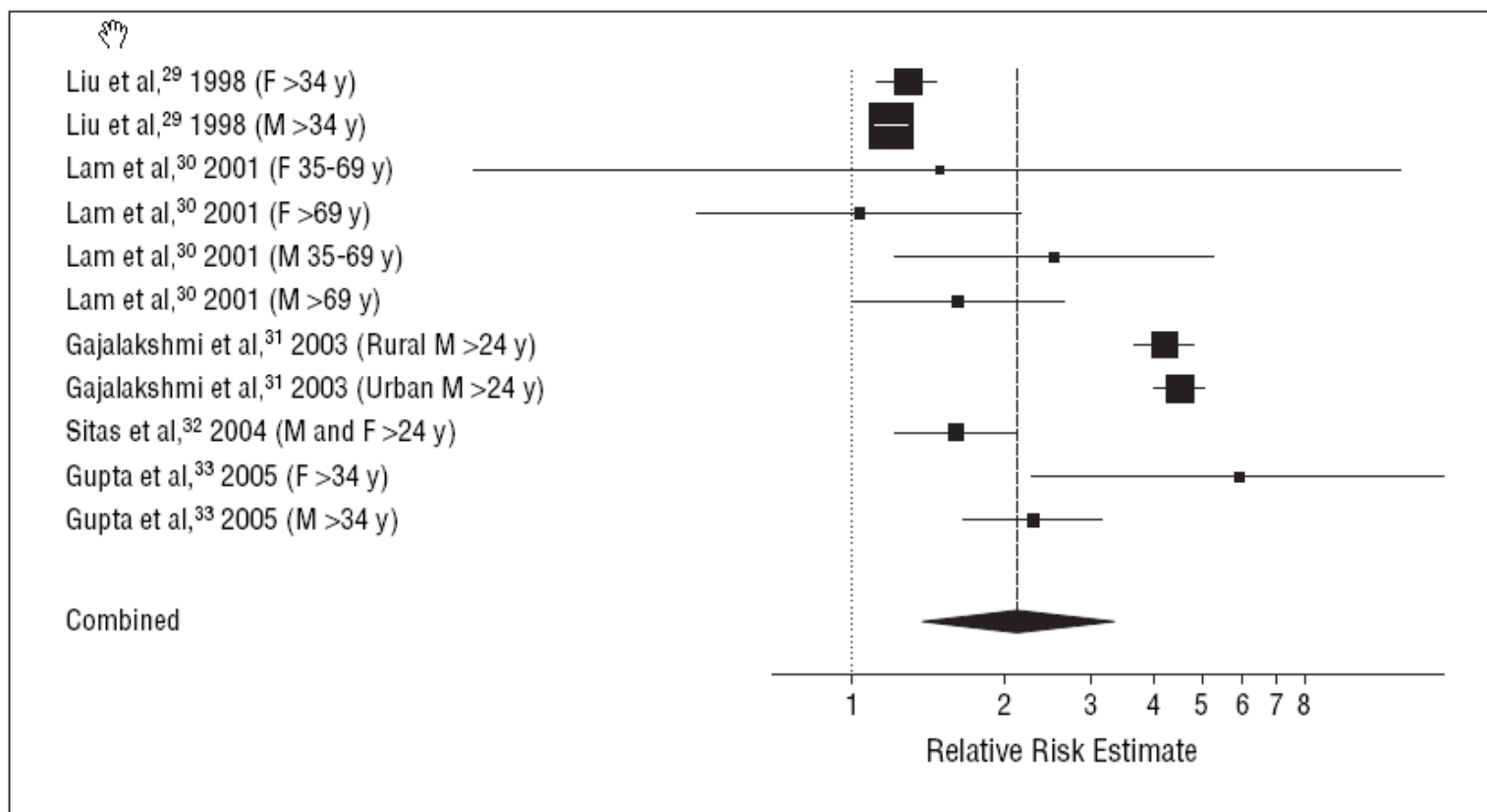
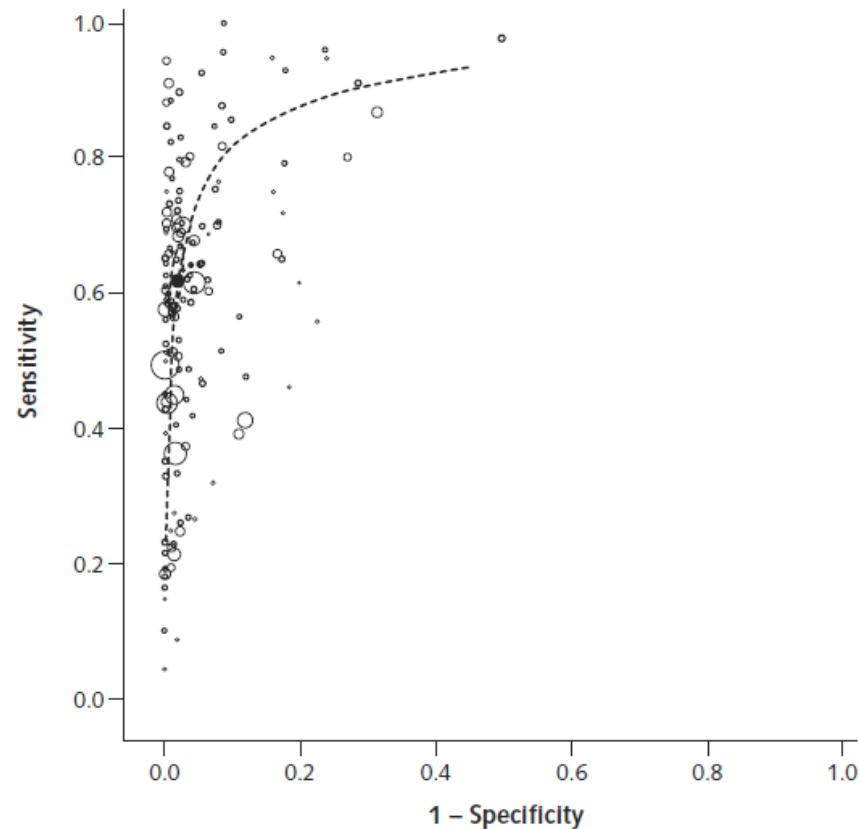


Figure 5. Forest plot of studies²⁹⁻³³ that examined smoking and tuberculosis mortality. The sex and age of the study population are shown on the y-axis.

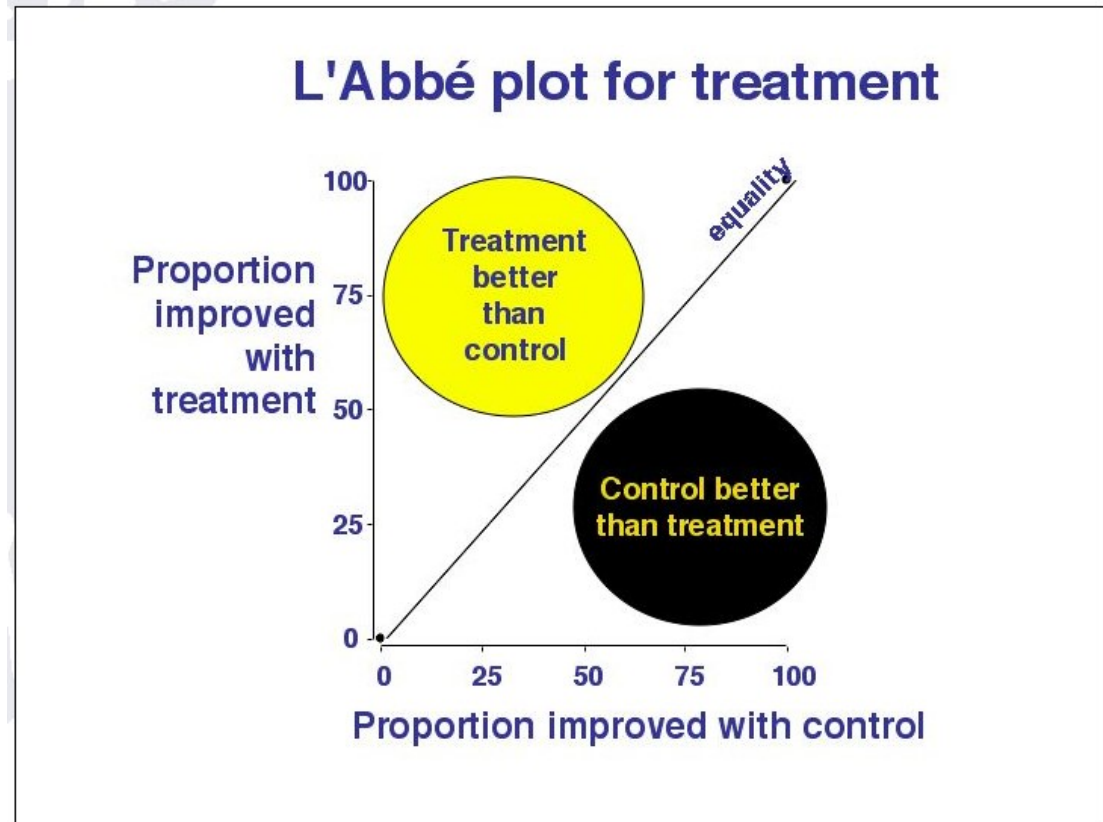
Figure 2. Hierarchical summary receiver-operating characteristic curve plot of rapid influenza diagnostic test studies.



Individual studies ($n = 159$) are shown as open circles whose size is proportionate to the size of the study. Summary point is shown as a closed circle, representing sensitivity estimates pooled by using bivariate random-effects regression model. The hierarchical summary receiver-operating characteristic curve is shown as a dashed line and is truncated outside the area for which data exist.

L'Abbe plot for heterogeneity

- Trials in which the experimental treatment proves better than the control ($EER > CER$) will be in the upper left of the plot, between the y axis and the line of equality (Figure). If experimental is no better than control then the point will fall on the line of equality ($EER = CER$), and if control is better than experimental then the point will be in the lower right of the plot, between the x axis and the line of equality ($EER < CER$).



3. Check for heterogeneity

- If significant heterogeneity is found:
 - Find out what factors might explain the heterogeneity
 - Can decide not to combine the data
- If no heterogeneity:
 - Can perform meta-analysis and generate a common, summary effect measure

Heterogeneity makes it hard to interpret pooled estimates

"We view the opposition of random-effects summaries and fixed-effects summaries as misleading and counterproductive, for the following reason: If the two summaries differ to a meaningful extent, there must be meaningful discrepancies (heterogeneity) among the study-specific effect estimates. In this situation, we contend that any summary will be inadequate."

Poole and Greenland (1999)

"[I]n drawing inferences from heterogeneous but logically related studies...the use of regression analysis to characterize differences in study outcomes may be more appropriate [than random-effects summarization]."

*DerSimonian and Laird (1985)**

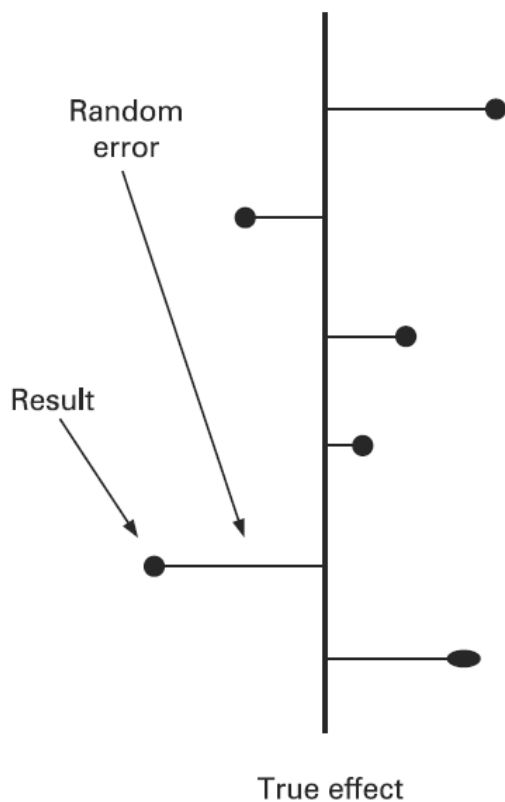
4. Perform meta-analysis

- Decide what data to combine
- Data types:
 - Continuous
 - Dichotomous
- Examples of measures that can be combined:
 - Risk ratio
 - Odds ratio
 - Risk difference
 - Effect size (Z statistic; standardized mean difference)
 - P-values
 - Correlation coefficient (R)
 - Sensitivity & Specificity of a diagnostic test

4. Perform meta-analysis

- Statistical models for combining data:
 - All methods are essentially compute weighted averages
 - Weighting factor is often the study size
 - Models:
 - Fixed effects model
 - Inverse-variance, Peto method, M-H method
 - Random effects model
 - DerSimonian & Laird method

Fixed-effect model



Random-effect model

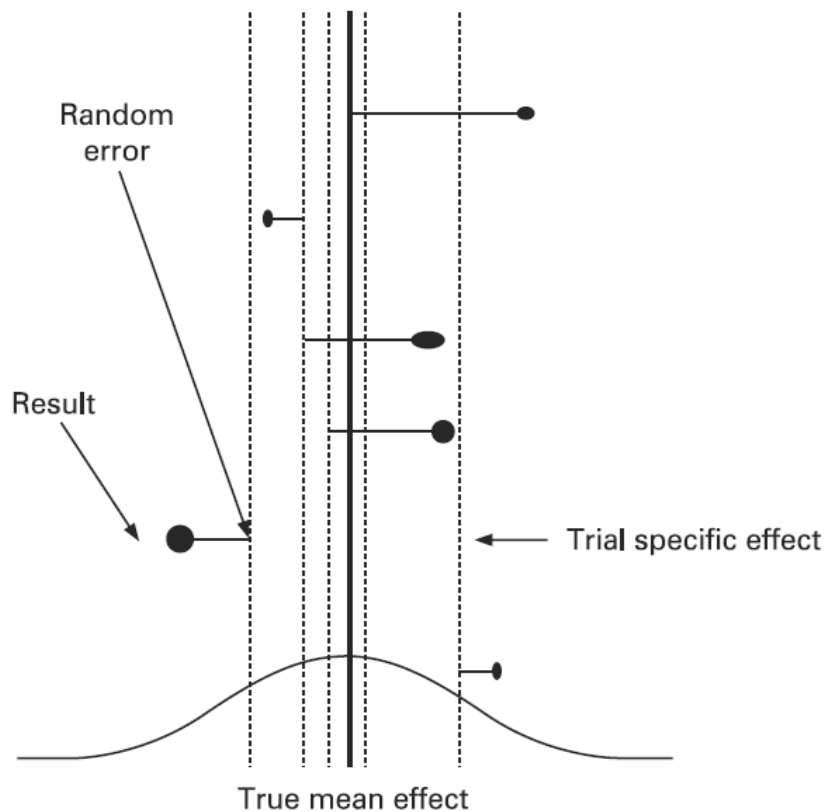
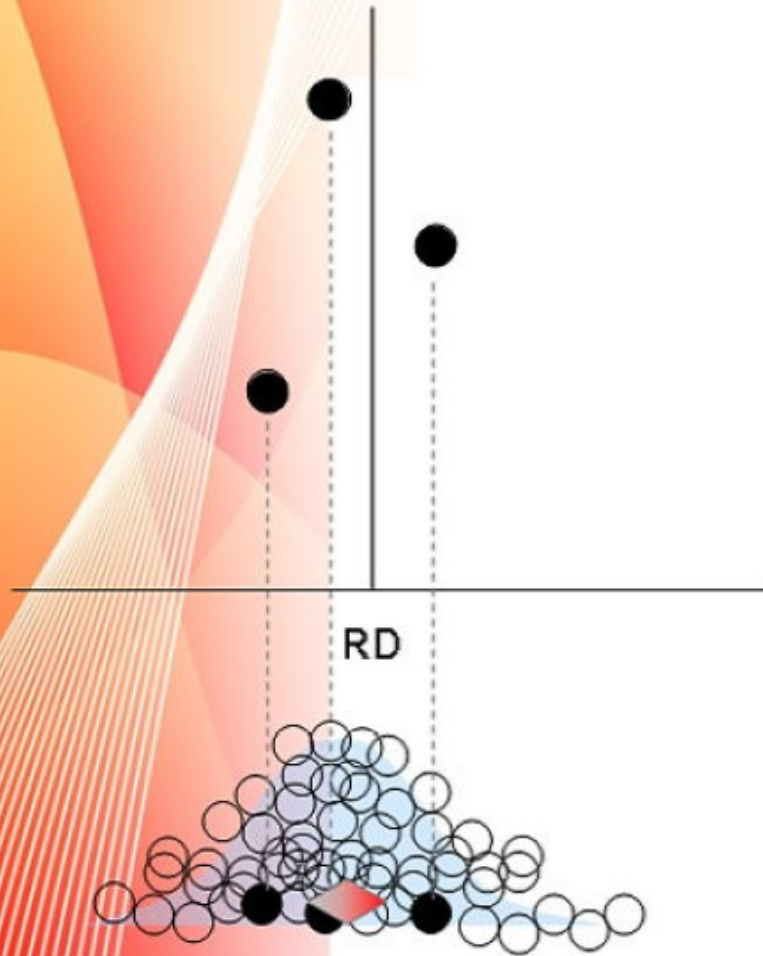
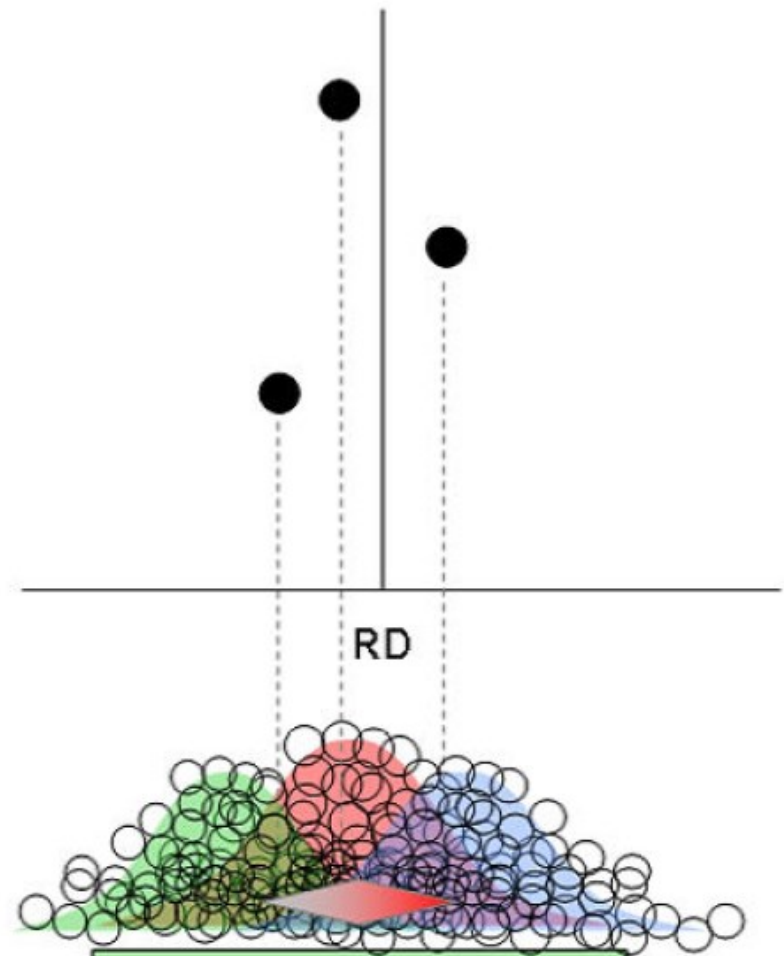


Figure 2. Graphical representation of the theoretical models used to combine data together.

Fixed and random effects models



Single, homogeneous source population

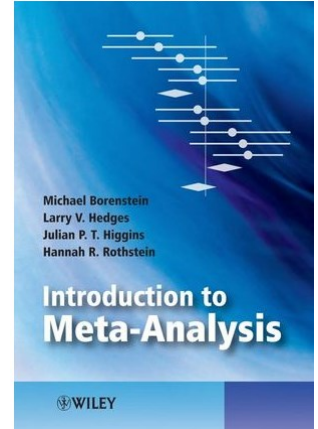


Multiple source populations

4. Perform meta-analysis

- Fixed effects model
 - based on the assumption that a single common (or 'fixed') effect underlies every study in the meta-analysis
 - For example, if we were doing a meta-analysis of ORs, we would assume that every study is estimating the same OR.
 - Under this assumption, if every study were infinitely large, every study would yield an identical result.
 - Same as assuming there is no statistical heterogeneity among the studies

	True effect	Observed effect
Study	●	■
Combined	▼	◆



Introduction to Meta-Analysis

WILEY

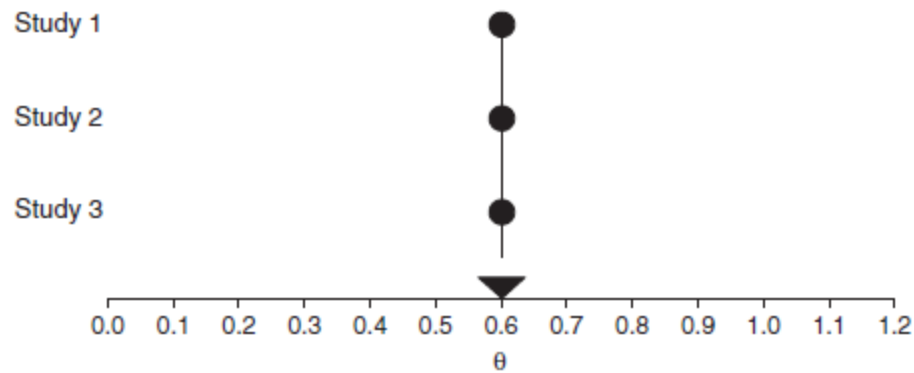


Figure 11.1 Fixed-effect model – true effects.

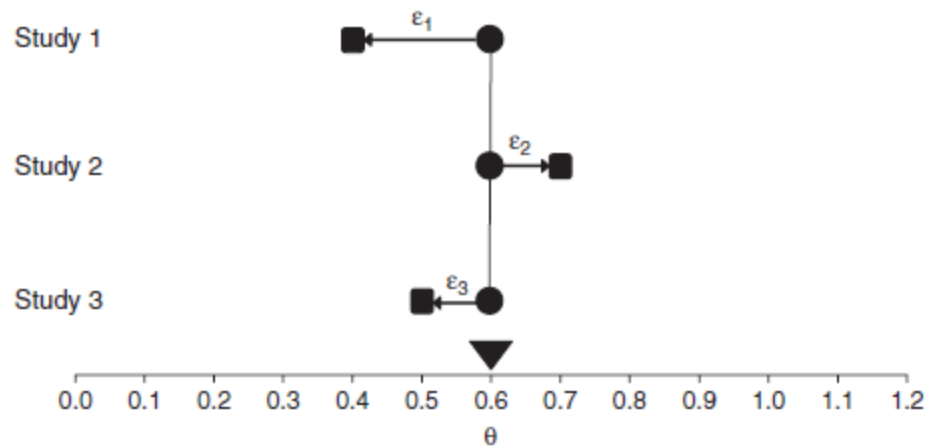


Figure 11.2 Fixed-effect model – true effects and sampling error.

Example of a fixed effects method (M-H)

Study 1

Disease

Treat ment		+	-
	+	a	b
	-	c	d

Study 2

Disease

Treat ment		+	-
	+	a	b
	-	c	d

$$OR_{MH} = \frac{\sum [a_i d_i / n_i]}{\sum [b_i c_i / n_i]}$$

Example of a fixed effects method (M-H)

Study 1: $n_1 = 200$

Disease

Treat ment	Disease	
	+	-
+	10	90
-	20	80

OR = 0.44

Study 2: $n_2 = 200$

Disease

Treat ment	Disease	
	+	-
+	12	88
-	16	84

OR = 0.72

$$OR_{MH} = \frac{\sum [a_i d_i / n_i]}{\sum [b_i c_i / n_i]} = (4 + 5.04) / (9 + 7.04) = OR_{MH} = 0.56$$

4. Perform meta-analysis

- Random effects model

- Makes the assumption that individual studies are estimating different true effects
 - we assume they have a distribution with some central value and some degree of variability
 - the idea of a random effects MA is to learn about this distribution of effects across different studies
- Random effects model:
 - Allows for random error plus inter-study variability
 - Results in wider confidence intervals (conservative)
 - Studies tend to be weighted more equally (relatively more weight is given to smaller studies)
 - Can be unpredictable (i.e. not stable)

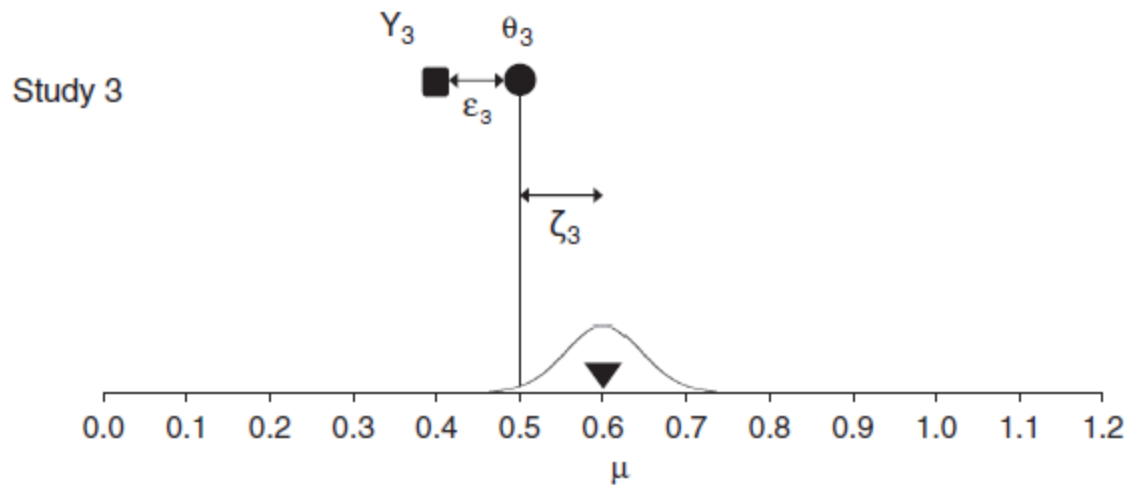


Figure 12.3 Random-effects model – true and observed effect in one study.

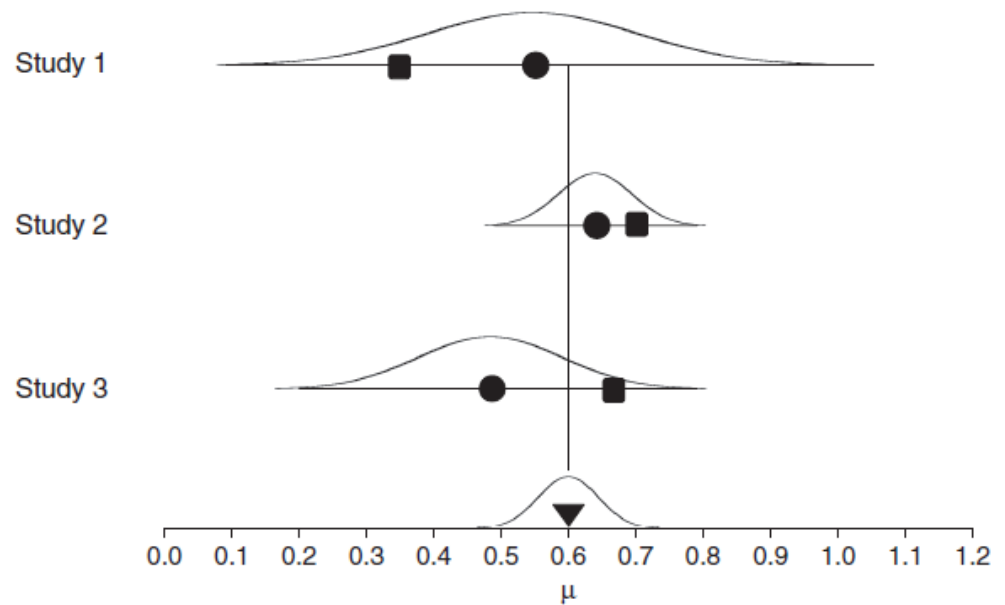


Figure 12.4 Random-effects model – between-study and within-study variance.

DerSimonian and Laird Model

$$Q = \sum_{i=1}^n w_i (y_i - \bar{y})^2,$$

where $w_i = \sigma_i^{-2}$, $\bar{y} = \sum_{i=1}^n w_i y_i / \sum_{i=1}^n w_i$ and n denotes the number of studies. Under the assumptions of the random effects model it can be shown that the expectation of Q is

$$E[Q] = (n-1) + \left(S_1 - \frac{S_2}{S_1} \right) \tau^2$$

where $S_r = \sum_{i=1}^n w_i^r$, which provides the DerSimonian and Laird estimate

$$\hat{\tau}_{DL}^2 = \max \left(0, \frac{Q - (n-1)}{S_1 - \frac{S_2}{S_1}} \right).$$

The corresponding estimate of treatment effect is

$$\hat{\mu}_{DL} = \frac{\sum_{i=1}^n \frac{y_i}{\sigma_i^2 + \hat{\tau}_{DL}^2}}{\sum_{i=1}^n \frac{1}{\sigma_i^2 + \hat{\tau}_{DL}^2}}.$$

RESEARCH METHODS & REPORTING

Interpretation of random effects meta-analyses

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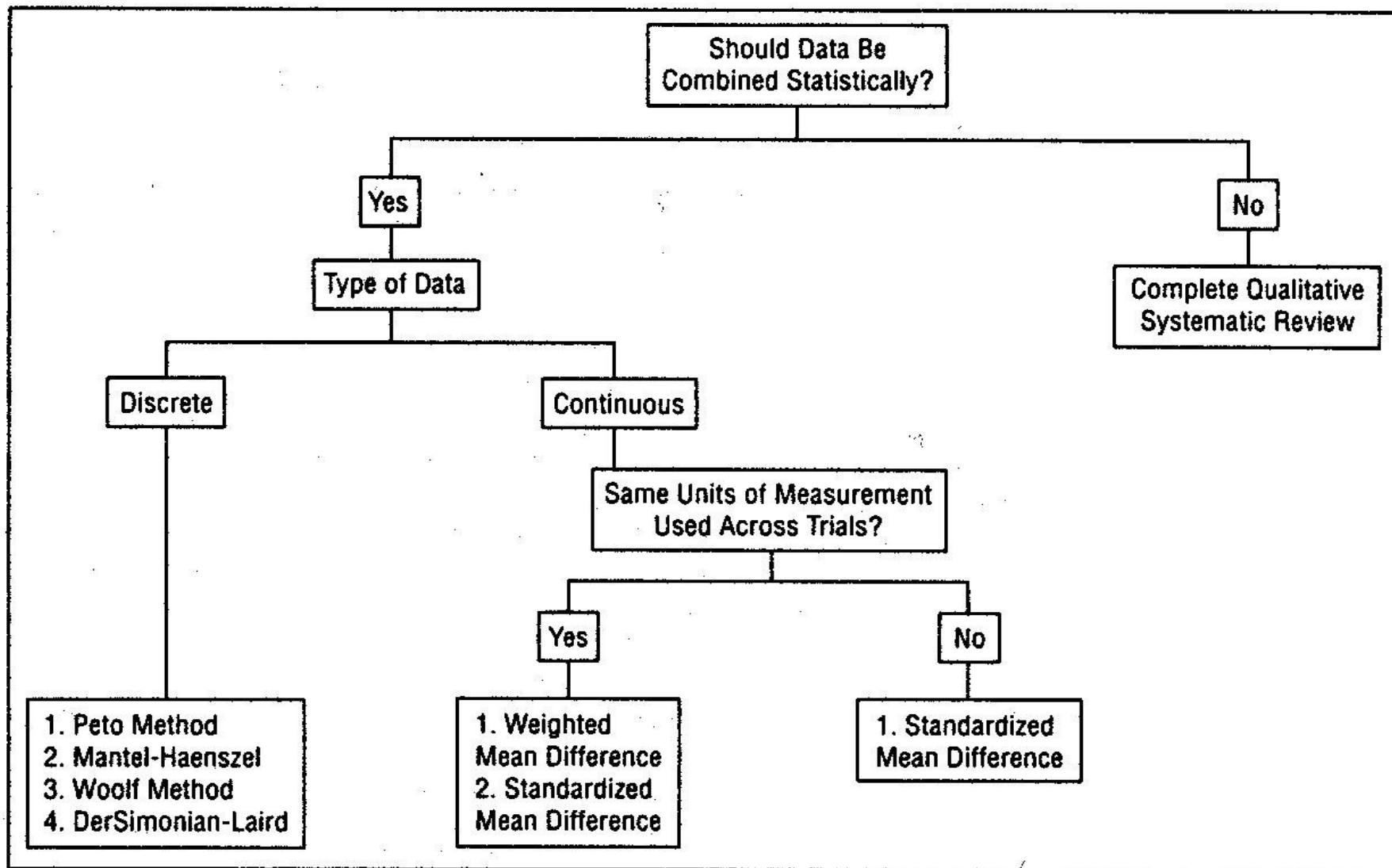
Accepted: 11 November 2010

Cite this as: *BMJ* 2011;342:d549
doi: 10.1136/bmj.d549

Summary estimates of treatment effect from random effects meta-analysis give only the average effect across all studies. Inclusion of prediction intervals, which estimate the likely effect in an individual setting, could make it easier to apply the results to clinical practice

mean difference in change in systolic blood pressure between the treatment group and the control group. Negative estimates indicate a greater blood pressure reduction for patients in the treatment group than the control group.

The two meta-analyses give identical summary estimates of treatment effect of -0.33 with a 95% confidence interval of -0.48 to -0.18 , but the first uses a fixed effect model and the second a random effects model. In the following two sections we explain why the summary result should be interpreted differently in these two examples because of the different meta-analysis models they use.



Algorithm of statistical choices available to systematic reviewers.

5. Identify factors that can explain heterogeneity

- If heterogeneity is found, use these approaches to identify factors that can explain it:
 - Graphical methods
 - Subgroup analysis
 - Sensitivity analysis
 - Meta-regression
- Of all these approaches, subgroup analysis is easily done and interpreted

Graphical exploration

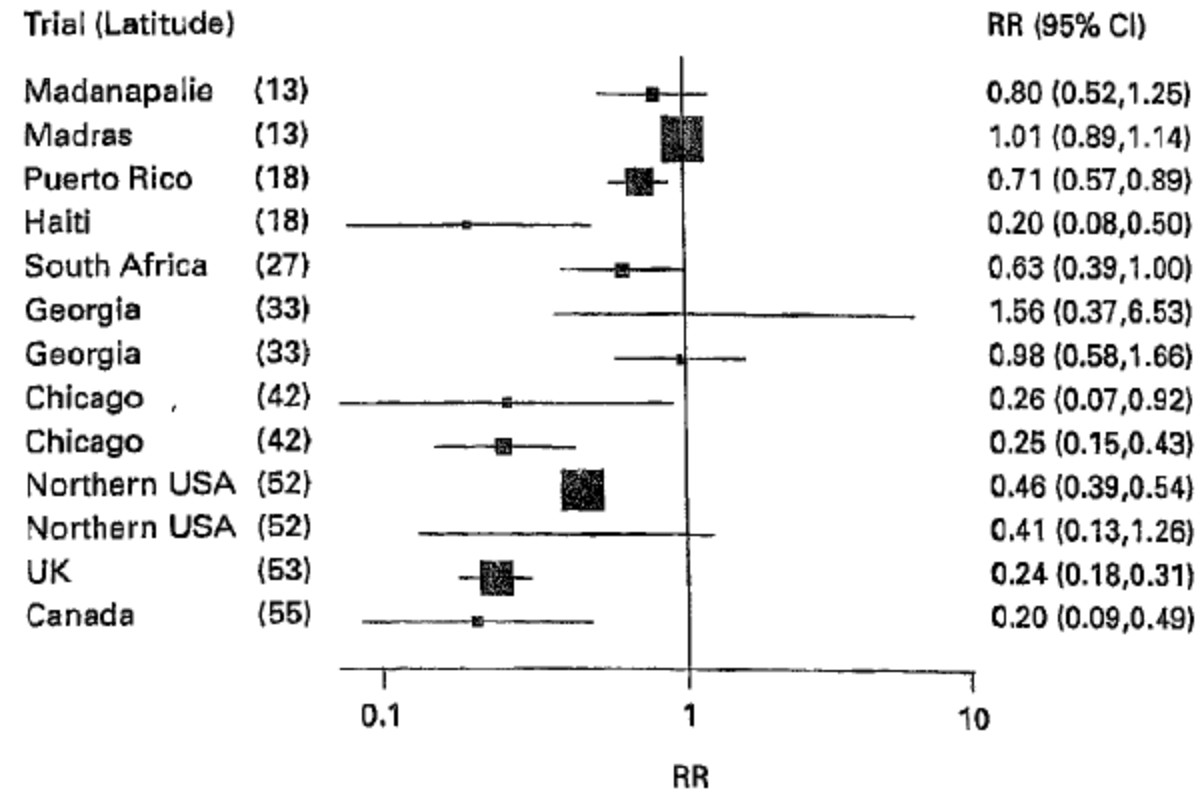


Fig 4. Forest plot of trials of BCG vaccine to prevent tuberculosis. Trials are ordered according to the latitude of the study location, expressed as degrees from the equator. No meta-analysis is shown (CI = confidence intervals, RR = relative risk) (adapted from Colditz *et al.*⁴⁷).

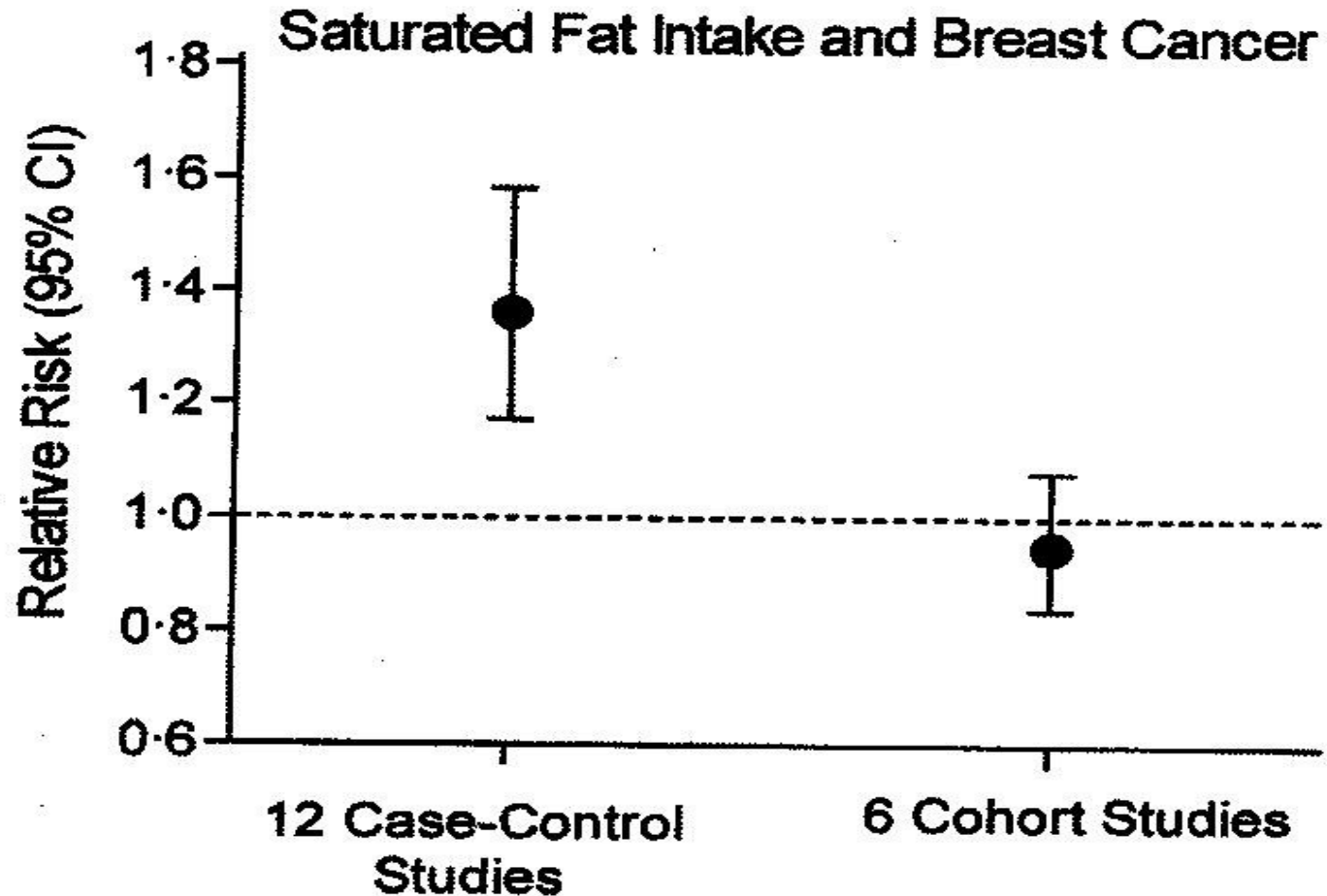
I-squared = 92%

Meta-analysis on efficacy of BCG vaccination for TB



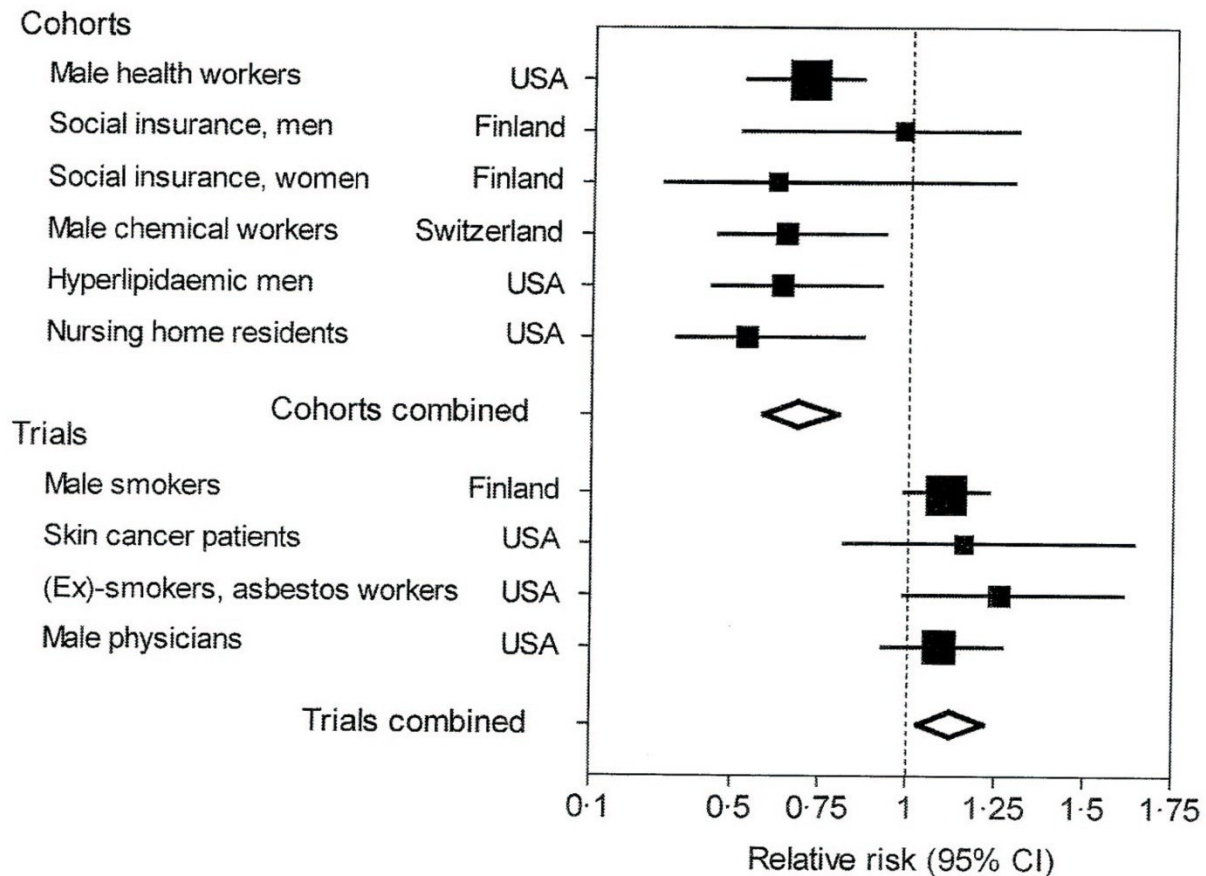
This photo is of a sign located on Interstate 89 in Vermont just south of the border with Quebec Province, Canada [source: Wikipedia]

Subgroup analysis: example



Subgroup analysis: example

Beta-carotene intake and cardiovascular mortality



Subgroup analysis: example

REVIEW

Annals of Internal Medicine

Accuracy of Rapid Influenza Diagnostic Tests

A Meta-analysis

Caroline Chartrand, MD, MSc; Mariska M.G. Leeflang, DVM, PhD; Jessica Minion, MD, MSc; Timothy Brewer, MD, MPH; and Madhukar Pai, MD, PhD

Table 2. Accuracy Estimates From Subgroup Analyses

Characteristic	Pooled Sensitivity (95% CI), %	P Value	Pooled Specificity (95% CI), %	P Value
Population				
Children (60 studies)	66.6 (61.6–71.7)	<0.001	98.2 (97.5–99.0)	0.135
Adults (33 studies)	53.9 (47.9–59.8)	Reference	98.6 (98.0–98.9)	Reference
Virus type				
Influenza A (72 studies)	64.6 (59.0–70.1)	0.62	99.1 (98.7–99.4)	<0.001
Influenza B (27 studies)	52.2 (45.0–59.3)	0.050	99.8 (99.7–99.9)	<0.001
Influenza A and B (47 studies)	62.3 (55.2–69.4)	Reference	96.1 (94.4–97.8)	Reference

“Considerable heterogeneity was found in the pooled estimates, as expected. Despite our attempts to explain it through the regression model, substantial heterogeneity remained unexplained.”

Exploring heterogeneity using meta-regression

- A meta-regression can be either a linear or logistic regression model
 - Can be weighted or unweighted
- Unit of analysis is a study (similar to an ecological study).
- Outcome variable: effect (e.g. log odds ratio)
- Covariates: study-level variables (e.g. Study quality, mean age of participants, etc)

Model: $\log \text{OR} = a + b_1X_1 + b_2X_2 + b_3X_3$

where, X_1, X_2 , etc are study level covariates

Exploring heterogeneity using meta-regression

- Limitations:

- Need sufficient data points (studies)
- Confounding is a concern
- False positives are likely and therefore need to pre-specify covariates (same as subgroup analysis)
- Need to limit the number of covariates (otherwise over-fitting is a problem)

STATISTICS IN MEDICINE
Statist. Med. 2002; **21**:1559–1573 (DOI: 10.1002/sim.1187)

How should meta-regression analyses be undertaken and interpreted?

Simon G. Thompson^{*†} and Julian P. T. Higgins

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6. Evaluate impact of study quality on results

- Narrative discussion of impact of quality on results
- Display study quality and results in a tabular format
- Weight the data by quality (not recommended)
- Subgroup analysis by quality
- Include quality as a covariate in meta-regression



"It's publish or perish, and he hasn't published."

7. Explore publication bias

- Studies with significant results are more likely
 - to be published
 - to be published in English
 - to be cited by others
 - to produce multiple publications
- Including only published studies can introduce publication bias
- Most reviews do not look for publication bias
- Methods for detecting publication bias:
 - Graphical: funnel plot asymmetry
 - Tests: Egger test, Rosenthal's Fail-safe N [all have low power]

Table 1. Steps in the Publishing Process Where Publication Bias May Intrude

Phases of research publication	Actions contributing to and/or resulting in publication bias
Preliminary and pilot studies	Small studies, more likely to be negative (discarded failed hypotheses), are unpublished—some under “industrial secret.”
Trial design, organization, and funding	Proposal selectively cites positive studies.
Institutional/ethics review board approval	No registries are kept of approved trials.
Study completion	Interim analysis shows that study is likely to be negative and project is dropped.
Report completion	Authors decide reporting a negative study is worthless and uninteresting, and no time or effort is assigned.
Report submission	Authors decide to forgo the submission of the negative study.
Journal selection	Authors decide to submit the report to a nonindexed, non-English-language, limited-circulation journal.
Editorial consideration	Editor decides that the negative study is not worth peer review process and rejects manuscript. If editor decides it is worth reviewing, manuscript goes to lower priority list.
Peer review	Reviewers conclude that the negative study does not contribute to the field and recommend rejection of the manuscript.
Author revision and resubmission	Author of rejected manuscript decides to forgo the submission of the negative study or to do it again at a later time to another journal (see “Journal selection”).
Report publication	Journal delays publication of the negative study.
Lay press report	The negative study is not considered newsworthy.
Electronic database indexing	Medline, EMBASE, Best Evidence do not scan or index articles in the journal/language of publication of the negative study.
Decision-maker retrieval	Health managers and policymakers do not retrieve the negative study to dictate policy.
Further trial evidence	New trial reports discuss their findings but do not cite the findings of the negative study.
Narrative review	Experts draft a review, but the negative study is never cited.
Systematic review	Reviewer goes to extremes to identify negative reports but misses the negative study. Industry-associated reviewer uses arbitrarily selected unpublished data “on file”; this further discredits incorporation of unpublished reports in systematic reviews.
Systematic review submission	Journal editors reject a meta-analysis because it included unpublished reports not exposed to the rigor of peer review. Review then follows the same path described here for the negative study.
Practice guidelines	Evidence-based guidelines are produced based on a systematic review that missed the negative study.
Funding opportunities	Further funding opportunities are identified without consideration of the negative study.

Systematic Review of the Empirical Evidence of Study Publication Bias and Outcome Reporting Bias

Kerry Dwan^{1*}, Douglas G. Altman², Juan A. Arnaiz³, Jill Bloom⁴, An-Wen Chan⁵, Eugenia Cronin⁶, Evelyn Decullier⁷, Philippa J. Easterbrook⁸, Erik Von Elm^{9,10}, Carrol Gamble¹, Davina Ghera¹¹, John P. A. Ioannidis^{12,13}, John Simes¹⁴, Paula R. Williamson¹

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Abstract

Background: The increased use of meta-analysis in systematic reviews of healthcare interventions has highlighted several types of bias that can arise during the completion of a randomised controlled trial. Study publication bias has been recognised as a potential threat to the validity of meta-analysis and can make the readily available evidence unreliable for decision making. Until recently, outcome reporting bias has received less attention.

Methodology/Principal Findings: We review and summarise the evidence from a series of cohort studies that have assessed study publication bias and outcome reporting bias in randomised controlled trials. Sixteen studies were eligible of which only two followed the cohort all the way through from protocol approval to information regarding publication of outcomes. Eleven of the studies investigated study publication bias and five investigated outcome reporting bias. Three studies have found that statistically significant outcomes had a higher odds of being fully reported compared to non-significant outcomes (range of odds ratios: 2.2 to 4.7). In comparing trial publications to protocols, we found that 40–62% of studies had at least one primary outcome that was changed, introduced, or omitted. We decided not to undertake meta-analysis due to the differences between studies.

Conclusions: Recent work provides direct empirical evidence for the existence of study publication bias and outcome reporting bias. There is strong evidence of an association between significant results and publication; studies that report positive or significant results are more likely to be published and outcomes that are statistically significant have higher odds of being fully reported. Publications have been found to be inconsistent with their protocols. Researchers need to be aware of the problems of both types of bias and efforts should be concentrated on improving the reporting of trials.

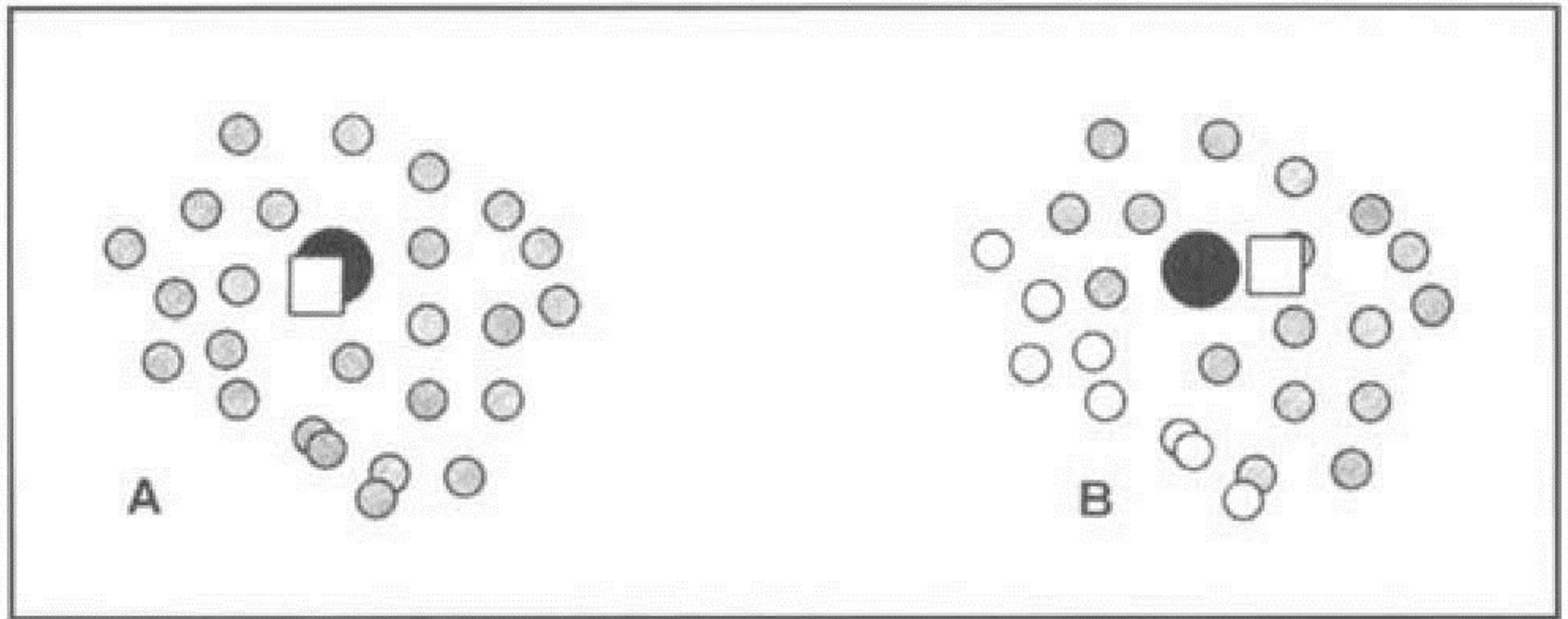
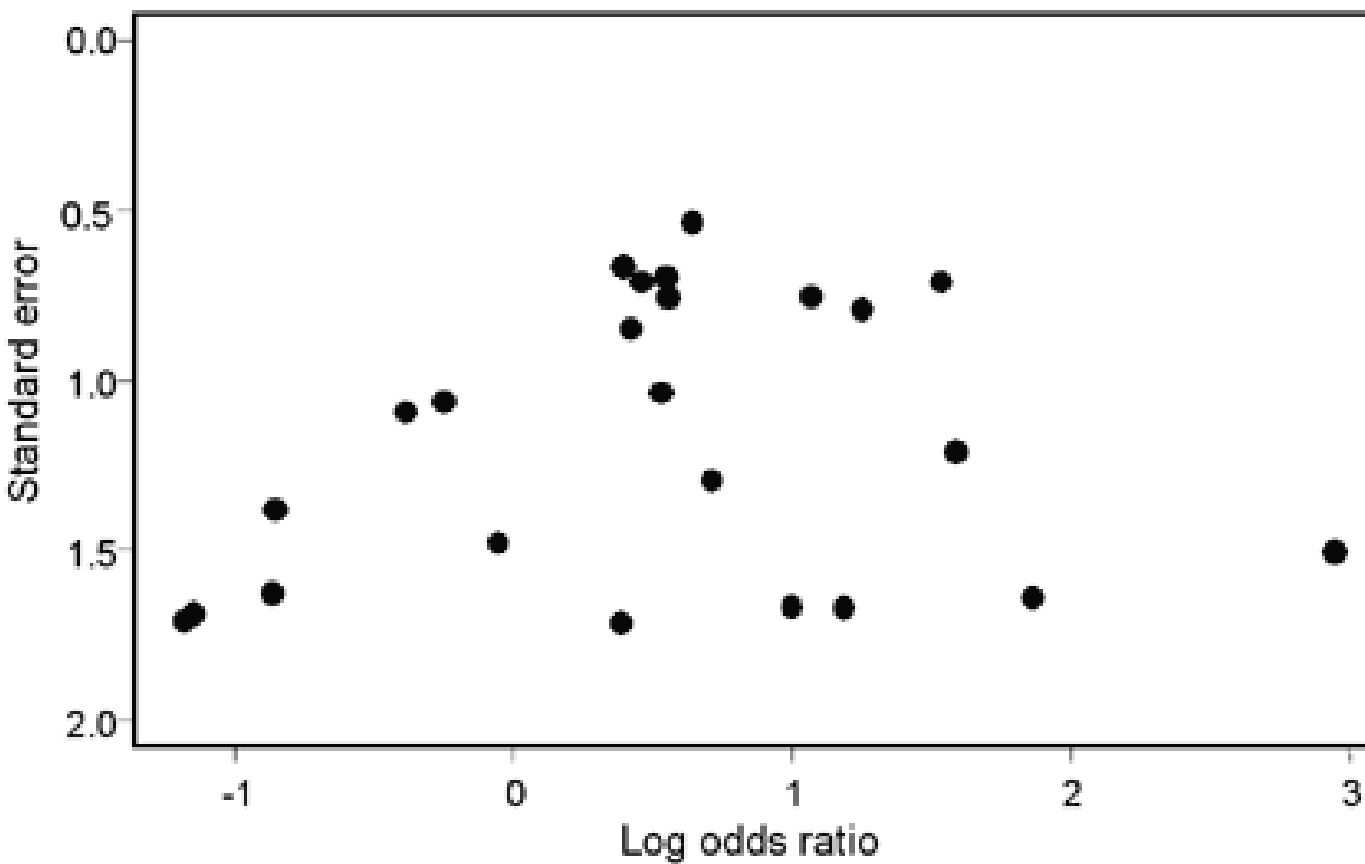
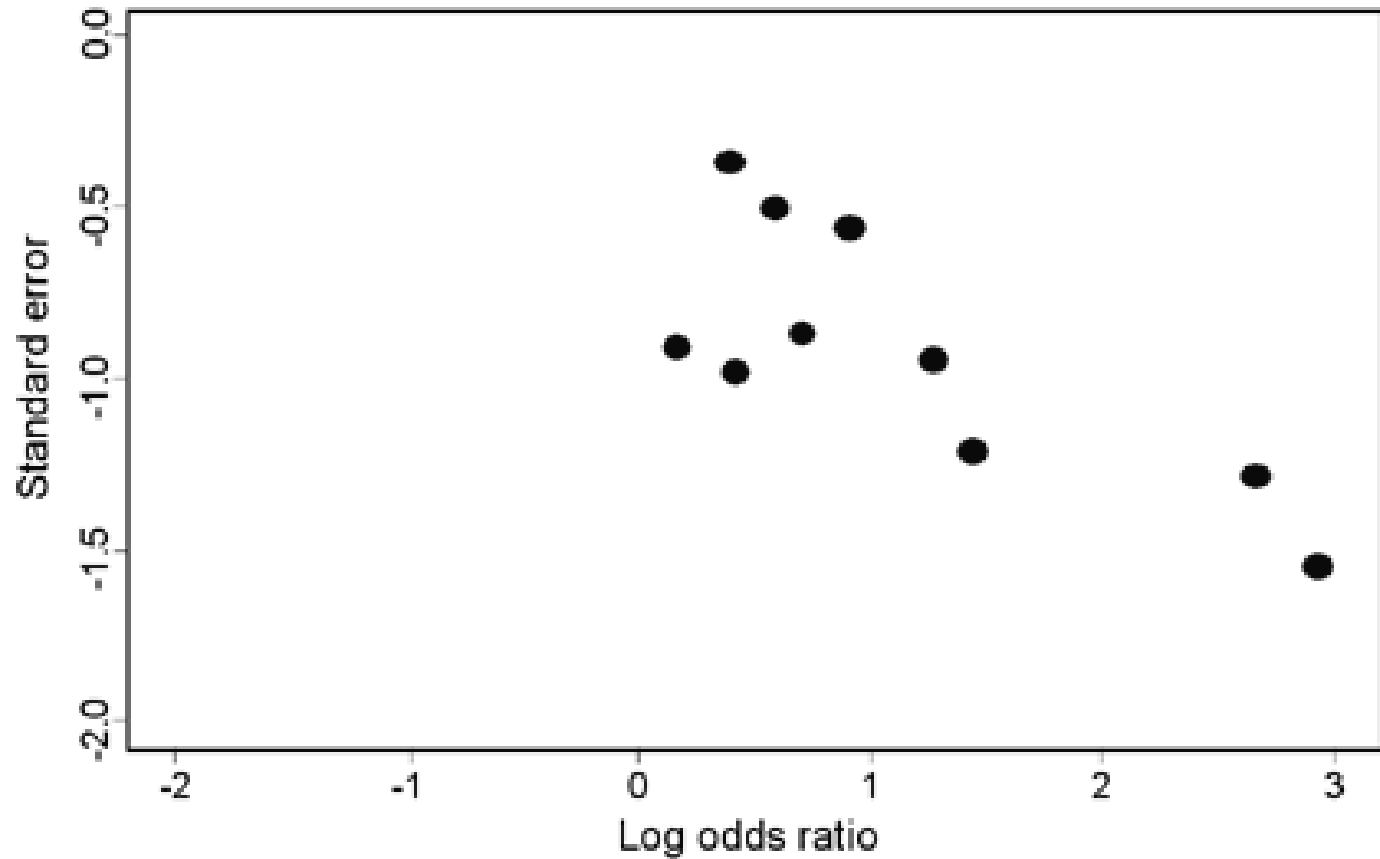


Figure 1. Publication bias. A, The black circle represents the underlying truth. The white square represents the pooled estimate from a systematic review of all the evidence (small shaded circles). B, The white circles represent evidence that was not identified by the reviewers because it was not published. Note the error in the pooled estimate (publication bias).



Funnel plot to detect publication bias



Testing for funnel plot asymmetry

Reference	Basis of test
(Begg 1994)	Rank correlation between standardized intervention effect and its standard error.
(Egger 1997a)	Linear regression of intervention effect estimate against its standard error, weighted by the inverse of the variance of the intervention effect estimate.
(Tang 2000)	Linear regression of intervention effect estimate on $1/\sqrt{N_{\text{tot}}}$, with weights N_{tot} .
(Macaskill 2001)*	Linear regression of intervention effect estimate on N_{tot} , with weights $S \times F / N_{\text{tot}}$.
(Deeks 2005)*	Linear regression of log odds ratio on $1/\sqrt{\text{ESS}}$ with weights ESS, where effective sample size $\text{ESS} = 4N_E \times N_C / N_{\text{tot}}$.
(Harbord 2006)*	Modified version of the test proposed by Egger et al., based on the 'score' ($O - E$) and 'score variance' (V) of the log odds ratio.
(Peters 2006)*	Linear regression of intervention effect estimate on $1/N_{\text{tot}}$, with weights $S \times F / N_{\text{tot}}$.
(Schwarzer 2007)*	Rank correlation test, using mean and variance of the non-central hypergeometric distribution.
(Rücker 2008)	Test based on arcsine transformation of observed risks, with explicit modelling of between-study heterogeneity.

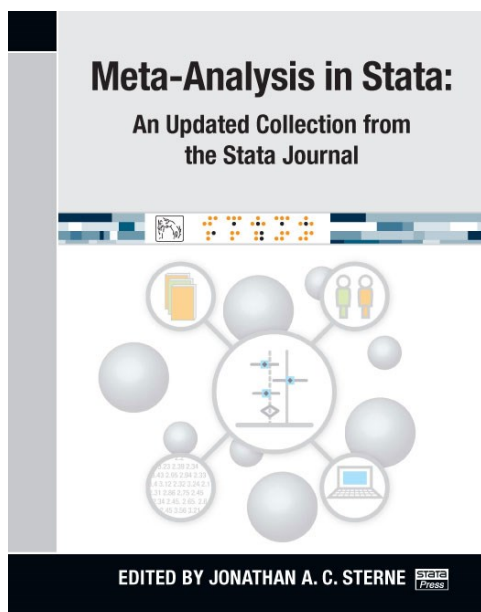
* Test formulated in terms of odds ratios, but may be applicable to other measures of intervention effect.

N_{tot} is the total sample size, N_E and N_C are the sizes of the experimental and control intervention groups, S is the total number of events across both groups and $F = N_{\text{tot}} - S$. Note that only the first three of these tests (Begg 1994, Egger 1997a, Tang 2000) can be used for continuous outcomes.

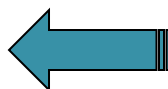
Recommendations by Cochrane

- As a rule of thumb, tests for funnel plot asymmetry should be used only when there are at least 10 studies included in the meta-analysis, because when there are fewer studies the power of the tests is too low to distinguish chance from real asymmetry.
- Tests for funnel plot asymmetry should not be used if all studies are of similar sizes (similar standard errors of intervention effect estimates).
- Results of tests for funnel plot asymmetry should be interpreted in the light of visual inspection of the funnel plot. For example, do small studies tend to lead to more or less beneficial intervention effect estimates? Are there studies with markedly different intervention effect estimates (outliers), or studies that are highly influential in the meta-analysis?
- When there is evidence of small-study effects, publication bias should be considered as only one of a number of possible explanations.
- Although funnel plots, and tests for funnel plot asymmetry, may alert review authors to a problem which needs considering, they do not provide a solution to this problem.
- Finally, review authors should remember that, because the tests typically have relatively low power, even when a test does not provide evidence of funnel plot asymmetry, bias (including publication bias) cannot be excluded.

Meta-analysis Software



- Free
 - RevMan 5 [Review Manager]
 - Meta-Analyst
 - Epi Meta
 - Easy MA
 - Meta-DiSc
 - Meta-Stat
- Commercial
 - Comprehensive Meta-analysis Version 2
 - MIX 2.0 Pro
 - Meta-Win
 - VEasy MA
- General stats packages (commercial)
 - Stata
 - SAS
 - R



Meta-analysis software

BMC Medical Research Methodology



Correspondence

Open Access

A systematic comparison of software dedicated to meta-analysis of causal studies

Leon Bax^{*1,2}, Ly-Mee Yu³, Noriaki Ikeda² and Karel GM Moons¹

Table 3: Meta-analysis software – basic feature comparison

	CMA	MetaAnalysis	MetaWin	MIX	RevMan	WEasyMA
General						
URL	meta-analysis.com	-	metawinsoft.com	mix-for-meta-analysis.info	cc-ims.net/RevMan	weasyrna.com
Corporate single user price	~\$1295.00	~\$75.00	~\$150.00	Free	\$650	~\$490.00
Student single user price	~\$395.00	~\$75.00	~\$75.00	Free	Free	~\$280.00
Download/program size	30 Mb	5 Mb	9 Mb	20 Mb/50 Mb	9 Mb	3 Mb
Compatibility	Windows	Windows	Windows	Windows	Windows	Windows
Last update	2006	2005	2002	2006	2005	2002
License	Single user	Single user	Single user	Open	Open	Single user
Input options						
Manual input	✓	✓	✓	✓	✓	✓
Copy & paste	✓		✓	✓	(✓)	
Text file import			✓	✓		
File import (Excel, other software)			✓	✓		
Descriptive dichotomous, e.g. n(total), n(y = 1)	✓	✓	✓	✓	✓	✓
Descriptive continuous, e.g. n, m, sd	✓		✓	✓	✓	
Comparative, e.g. theta, se/var	✓		✓	✓	✓	
Multi-format (mixed in one data set)	✓					
Single data input/selection	✓	✓		✓	✓	✓
Maximum number of studies	Unlimited	Unlimited	Unlimited	100	Unlimited	Unlimited
Information sources						
Within-program HTML help			✓	(✓)	✓	(✓)
Printable manual	✓	✓	✓		✓	
Description of methods/calculations		(✓)	✓	(✓)	✓	
Additional information sources (PDFs/tutorials)	✓		✓	✓		
Up-to-date website	✓		✓	✓	✓	x

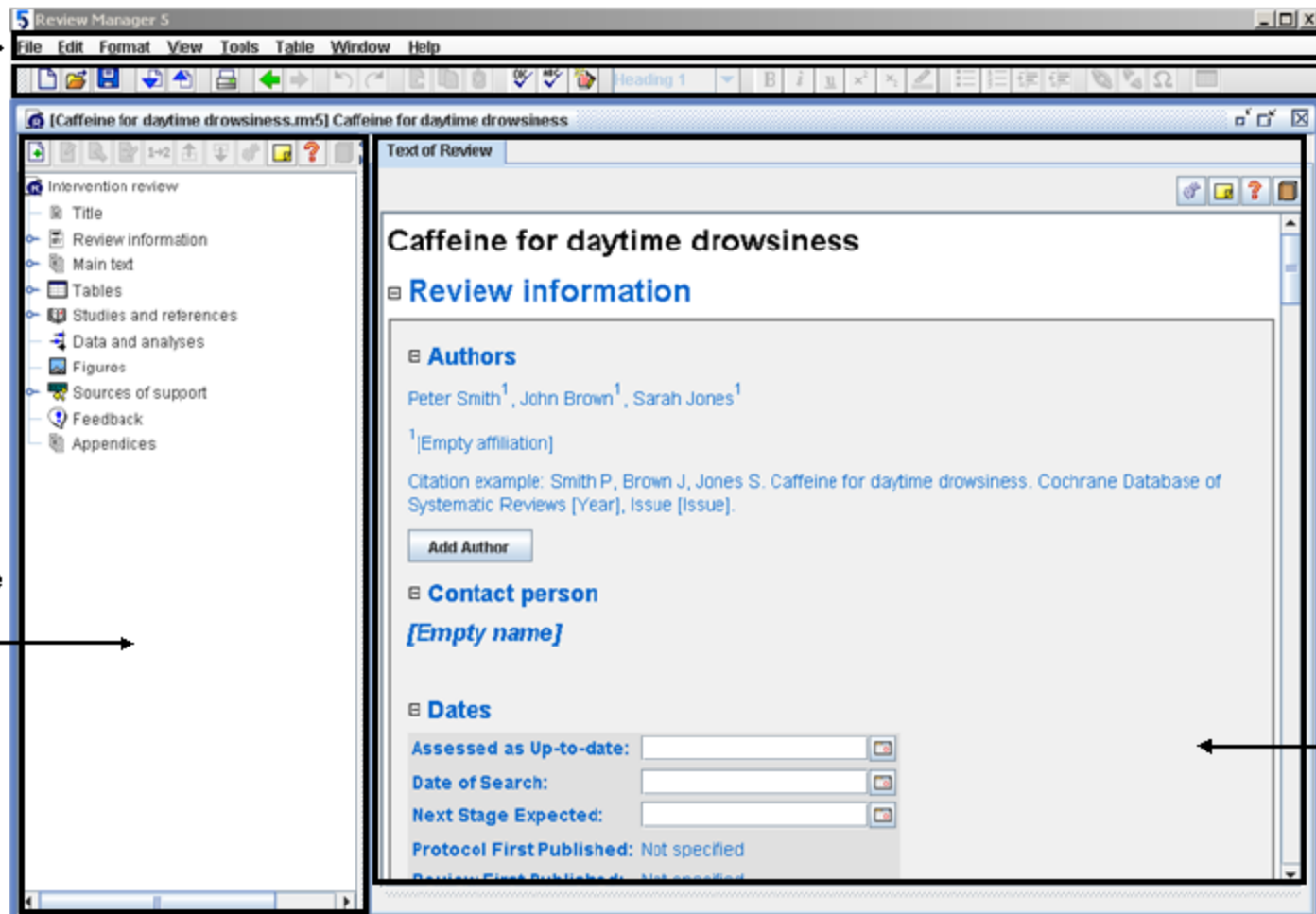
RevMan 5

Menu
Bar

Toolbar

Outline
Pane

Content
Pane





Review

```

1 use "C:\Documents and S
2 metan deaths1 deaths0
3 metan deaths1 deaths0
4 metan deaths1 alive1 de

```

```

16 | 1.096 0.903 1.331 2.34
17 | 0.889 0.456 1.734 0.25
18 | 0.893 0.644 1.239 1.13
19 | 0.611 0.435 0.858 1.66
20 | 0.931 0.761 1.139 2.88
21 | 0.894 0.841 0.950 32.99
22 | 0.856 0.811 0.904 44.01

```

```

-----
M-H pooled RR | 0.874 0.844 0.906 100.00
-----

```

```

Heterogeneity chi-squared = 34.6
I-squared (variation in RR attrib

```

```

Test of RR=1 : z= 7.41 p = 0.00
option plot not allowed
r(198);

```

```

. metan deaths1 alive1 deaths0 alive1

```

```

-----
Study | OR [95% CI] | % Weight
-----
Fletcher | 0.159 | 0.18
Dewar | 0.471 | 0.27
1st European | 1.460 | 0.54
Heikinheimo | 1.248 | 0.75
Italian | 1.012 | 0.78
2nd European | 0.635 | 3.76
2nd Frankfurt | 0.378 | 1.20
1st Australian | 0.754 | 1.41
NHLBI SMIT | 2.587 | 0.36
Valere | 1.061 | 0.26
Frank | 0.959 | 2.13
UK collab | 0.876 | 0.04
Klein | 3.200 | 2.70
Austrian | 0.562 | 0.14
Lasierra | 0.222 | 1.88
N German | 1.215 | 0.22
Witchitz | 0.778 | 1.13
2nd Australian | 0.806 | 2.00
3rd European | 0.416 | 2.80
ISAM | 0.872 | 32.49
GISSI-1 | 0.807 | 44.82
ISIS-2 | 0.746 | 100.00
-----
M-H pooled OR | 0.774
-----

```

```

Heterogeneity chi-squared = 31.5
I-squared (variation in OR attrib

```

```

Test of OR=1 : z= 7.76 p = 0.00

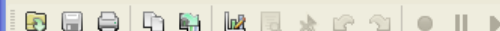
```

Variables

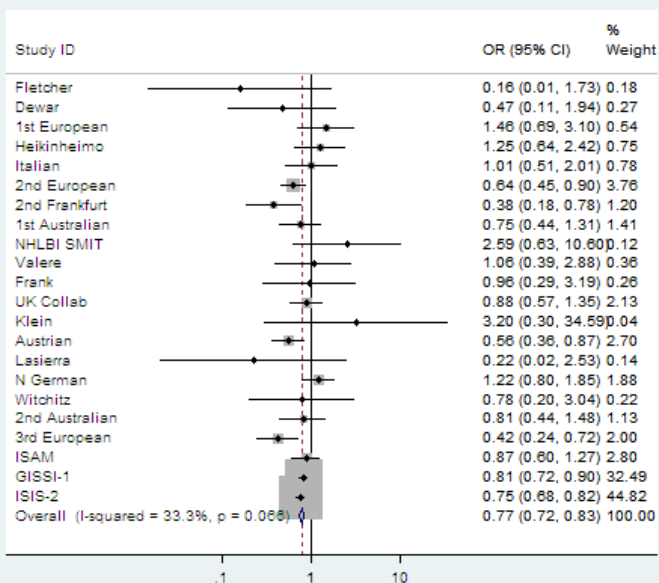
Name	Label
trial	Trial number
trialnam	Trial name
year	Year of publication
deaths1	Treated deaths
deaths0	Control deaths
alive1	
alive0	
logor	
selogor	
_SS	Sample size
_ES	OR
_selogES	se(logOR)
_LCI	Lower CI (OR)
_UCI	Upper CI (OR)
_WT	M-H weight

Stata Graph - Graph

File Edit Object Graph Tools Help



Graph



Command

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META-ANALYSIS MADE EASY WITH MIX 2.0

MIX 2.0 is a statistical add-in for performing meta-analysis in Excel. The free Lite version is for educational purposes and contains datasets that are used in the most authoritative books on meta-analysis. The paid Pro version has additional features that enable users to analyze their own datasets. For more information, visit the [About page](#).

TOP RATING ON CNET DOWNLOAD.COM

MIX 2.0 Lite was reviewed by editors of CNET Download.com and received the highest rating of 5 stars!

CNET Editors' Rating:
★★★★★
Spectacular

'This sophisticated tool ... definitely makes it easy to apply meta-analysis to Excel.'
'... we highly recommend this free add-on.'

To read the full review, [click here](#).

DOWNLOAD MILESTONES

As 2013 progresses, MIX 2.0 continues to be one of the most popular software programs for meta-analysis. The Pro version has been downloaded 7,000+ times from CNET Download.com, while the Lite version has passed the 10,000-download mark! Both versions are available for download from the sites listed on the [Download page](#).

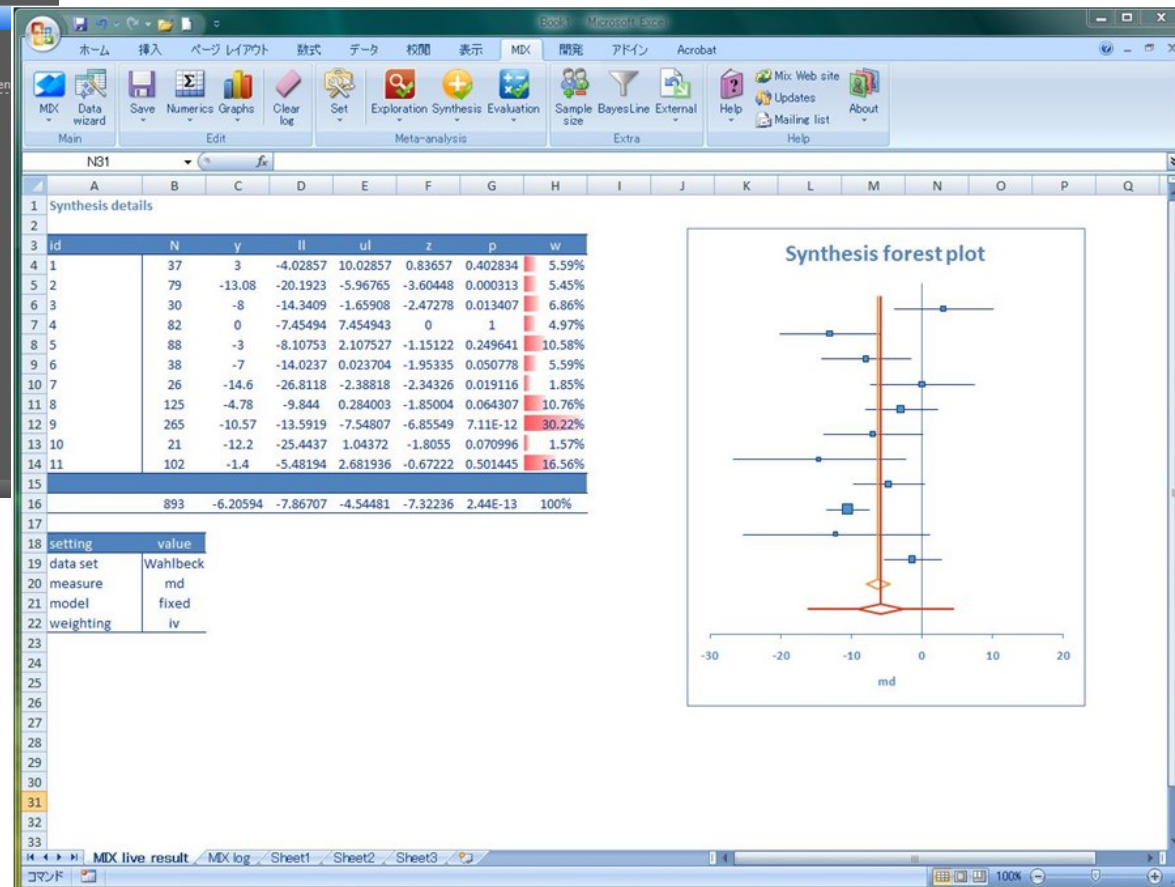
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COMPREHENSIVE META-ANALYSIS

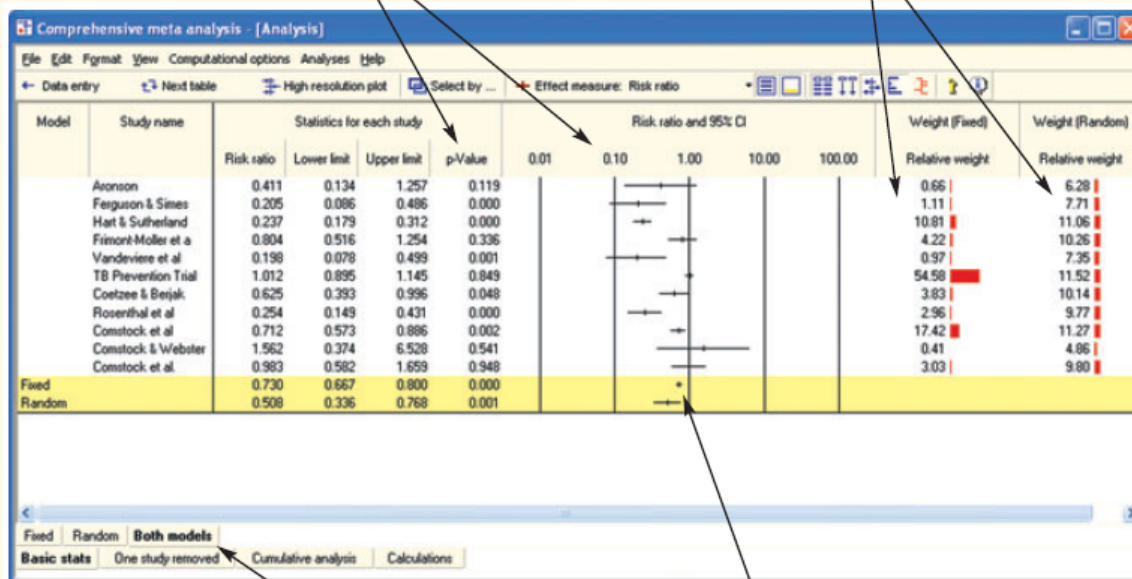
A computer program for meta-analysis

Version 2

<http://www.meta-analysis.com/>

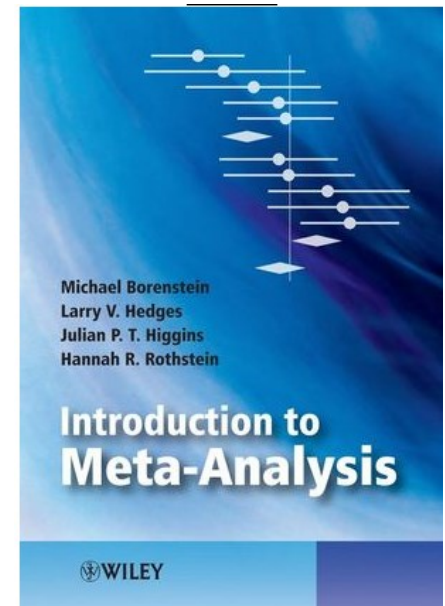
The program shows the effect size and confidence interval for each study.

The program shows the relative weight assigned to each study using fixed and random effects.



Select the computational model.

The program shows the combined effect size and confidence interval using fixed and random effects.

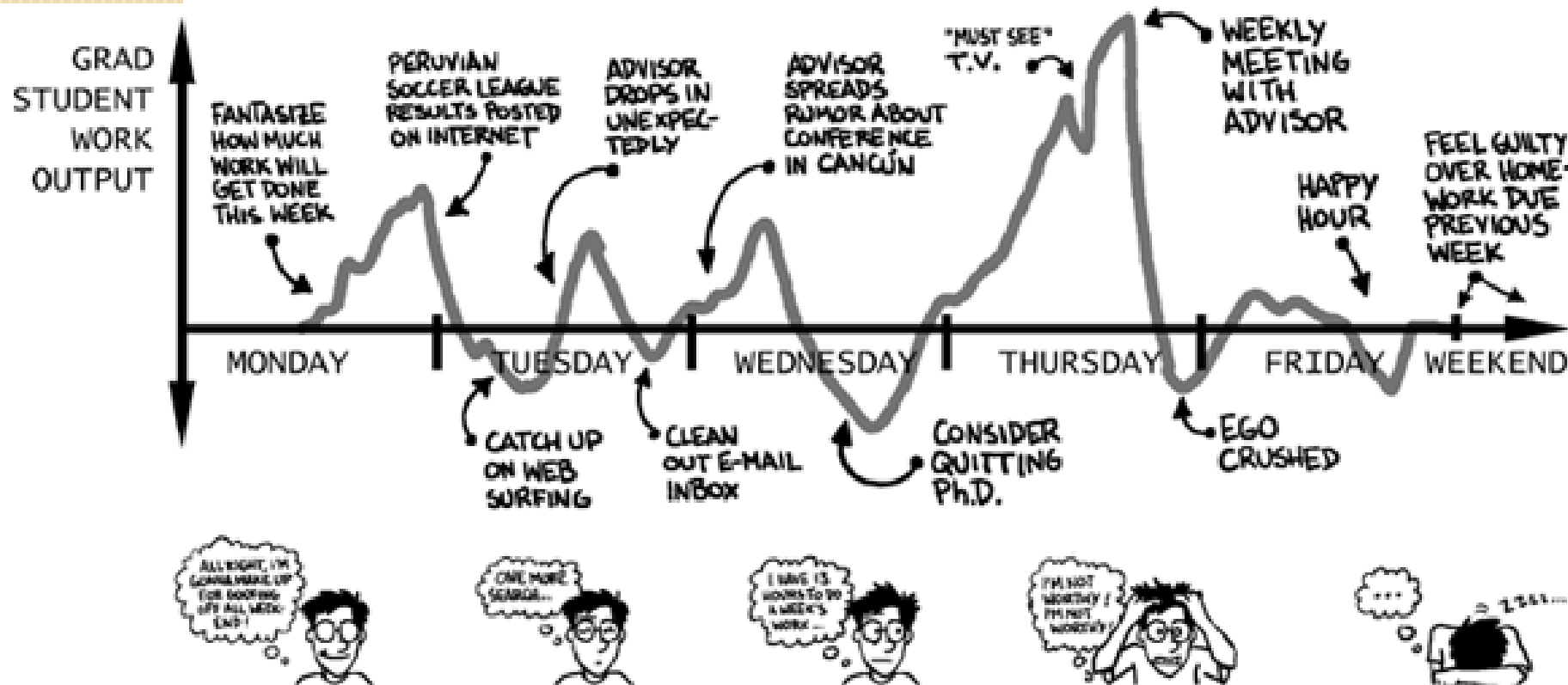


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