

Analytical Methods: Summary measures

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Analytical Methods

- Always start the meta-analysis with a “visual meta-analysis” (i.e., a great table 1 and forest plot).
 - A clinician should be able to interpret the results
- Step 1: Calculate a summary measure = “weighted mean effect estimate”
 - You can combine anything, but use judgment
- Step 2: Assess for heterogeneity
 - Heterogeneity is not always a problem
- Step 3: Assess for publication bias
 - Both visual and statistical methods
- Step 4: Perform subgroup/sensitivity analyses
 - Ideally specify these a priori

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How do you create a summary measure?

- Clinical example: 5 year old girl presents with ear pain and is found to have an acute otitis media.
- Should she get antibiotics?

Research Questions:

1. In children with OM, are antibiotics effective for pain relief?
2. In children with OM, do antibiotics reduce the rate of complications (mastoiditis, hearing problems)?

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3 studies are identified (examining effect of Abx on Pain)

- Study 1: N = 100 RR=1.41
- Study 2: N=200 RR=0.98
- Study 3: N=300 RR=1.01
- You could take the average effect: $(1.41 + 0.98 + 1.01) / 3 = 1.13$
- Is this a good summary measure?

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Summary measure weighted by sample size

- Provide “weight” for studies based on their sample size

Study	N	RR
1	100	1.41
2	200	0.98
3	300	1.01
Total	600	

$$\text{summary effect estimate} = \frac{\sum (N_i \times \text{effect estimate}_i)}{\sum (N_i)} = \frac{640}{600} = 1.07$$

More refined: Provide “weight” by using inverse of variance

Study	N	RR	Var RR	Weight
1	100	1.41	3.0	0.33
2	200	0.98	0.1	10
3	300	1.01	0.05	20
Total	700			

$$\text{Summary effect estimate} = \frac{\sum (\text{weight}_i \times \text{effect estimate}_i)}{\sum (\text{weight}_i)} = \frac{30.5}{30.3} = 1.00$$

The summary measure

- To perform a meta-analysis we compute an **effect size** and **variance** for each study, and then compute a **weighted mean** of these effect sizes.
- Odds ratios, rate ratio, risk ratios, risk differences, standardized mean differences or correlations are the measure of association in various study design.

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What is an Effect Size?

- Effect size – a way of expressing results in a common metric
- Units – standard deviation

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Table 3.1 Roadmap of formulas in subsequent chapters.

Effect sizes based on means (Chapter 4)
Raw (unstandardized) mean difference (<i>D</i>)
Based on studies with independent groups
Based on studies with matched groups or pre-post designs
Standardized mean difference (<i>d</i> or <i>g</i>)
Based on studies with independent groups
Based on studies with matched groups or pre-post designs
Response ratios (<i>R</i>)
Based on studies with independent groups
Effect sizes based on binary data (Chapter 5)
Risk ratio (<i>RR</i>)
Based on studies with independent groups
Odds ratio (<i>OR</i>)
Based on studies with independent groups
Risk difference (<i>RD</i>)
Based on studies with independent groups
Effect sizes based on correlational data (Chapter 6)
Correlation (<i>r</i>)
Based on studies with one group

- Results extracted from study reports may need to be **converted** to a consistent, or usable, format for analysis .

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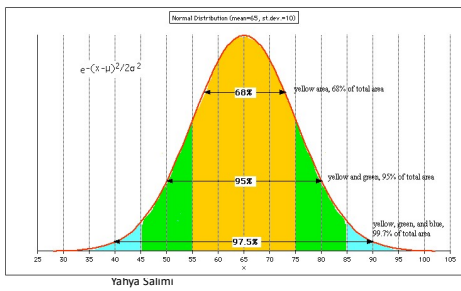
Effect Size

- ES =
$$\frac{X_1 - X_2}{SD_{pooled}}$$

1. ES increases as difference between means increases
2. ES increases as SD decreases
3. ES is expressed in units of SD
4. Summary ES combines the weighted ES from each study.

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Effect Size




Analytical model

- To compute the **weighted mean** we generally assign **more weight** to the **more precise** studies, but the rules for assigning weights depend on our **assumptions about** the distribution of true effects.

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Fixed-effects model

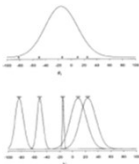


Fixed-effects meta-analysis assumes that the intervention has a single true effect.

$$\frac{\sum_{i=1}^k W_i Y_i}{\sum_{i=1}^k W_i}$$

$$W_i = \frac{1}{S_i^2}$$

Random-effects model



Random-effects meta-analysis assumes that the effect of the intervention varies across studies.

$$W_i(\tau) = \frac{1}{S_i^2 + \tau^2}$$

$$\frac{\sum_{i=1}^k W_i(\tau) Y_i}{\sum_{i=1}^k W_i(\tau)}$$

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Fixed-effect model

- Under the fixed-effect model we assume that there is **one true effect** size that underlies **all the studies** in the analysis, and that all **differences** in observed effects are due to **sampling error**.

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Random-effects model

- Under the **random-effects model** we allow that the true effect size might **differ from study to study**.
- The term "Random" reflects the fact that the studies included in the analysis are assumed to be a **random sample of all possible studies** that **meet the inclusion criteria** for the review.
- For example, the effect size might be higher (or lower) in studies where the participants are older, or more educated, or healthier than in other studies, or when a more intensive variant of an intervention is used.

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Fixed-effects model OR Random-effects model

Which should I choose?

- Good news – most software produces both
- Both theoretical and practical reasons for choosing
 - Generalizability
 - What is the focus of this meta-analysis?
 - Can I assume this is the whole population?

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The mistake to avoid

- Some researchers start the analysis by selecting the fixed-effect model. They then test perform a statistical test for heterogeneity in effect sizes (the Q-test).
- If the test for heterogeneity is not statistically significant, they conclude that the fixed-effect model is consistent with the data, and use this model in the analysis.
- If the test for heterogeneity is statistically significant they conclude that the fixed-effect model is not consistent with the data, and use the random-effects model in the analysis.

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Why does it matter which model we use?

- If we should be using the random-effects model and (by mistake) employ the fixed-effect model, then it's likely that:
 - The **estimate of the mean** will be **incorrect**
 - The **standard error** will be **incorrect**
 - The **test of significance** for the mean will be **incorrect**
 - The **confidence interval** about the mean effect will be **too narrow**

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In sum

- The selection of the correct statistical model is critically important.
- We should choose the model that fits the sampling frame.
- We should not choose a model based on the statistical test for heterogeneity

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Fixed-effect model

- Based on the fixed-effect model, there are several methods to estimate the weighted mean (or pooled) effect size.
- -Inverse-variance weighted estimation method (i.e. inverse of each study's variance)
- -Peto method
- -Mantel-Haenszel method
- -Maximum likelihood estimation method

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Fixed-effect model

	Yes	No	Total
Treatment	a1	b1	n1
Control	a2	b2	n2

IVW OR (i.e. ES): $(a1*b2)/(b1*a2)$ Weight: $1/(1/a1+1/b1+1/a2+1/b2)$

M-H OR (i.e. ES): $(a1*b2)/(b1*a2)$ Weight: $(b1+a2)/(a1+b1+a2+b2)$

Peto OR (i.e. ES): $\frac{observed_i - expected_i}{I_i}$
 $I_i = \frac{(a1_i + b1_i)^2 * (a2_i + b2_i)^2 * (a1_i + a2_i) * (b1_i + b2_i)}{n_i^2 * (n_i - 1)}$
 Weight: I_i

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Random-effects model

- Based on the random-effects model, methods to estimate the weighted mean ES (i.e. based on how to calculate tau-squared)
- -Weighted least squares estimation method (called DerSimonian-Laired (DL) method)
- -Maximum likelihood estimation method
- -Restricted maximum likelihood estimation method

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Example

- Antibiotics for acute bronchitis.
- After search and application of inclusion/exclusion criteria, 8 studies were included.

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RCTs in Acute Bronchitis

Study, yr	N	Abx	Outcome	Result*
Stott, 76	207	Doxy	Days of Yellow Spit	0.6 (-0.2 to 1.4)
Franks, 84	54	TMP/S	Cough Amount Score	0.2 (-0.2 to 0.6)
Williamson, 84	69	Doxy	Days of Purulent Sputum	-0.2 (-1.2 to 0.8)
Dunlay, 87	45	Erythro	Sputum production score	0.5 (0.1 to 0.9)
Scherl, 87	31	Doxy	Days of sputum	1.9 (-0.2 to 4.0)
Verheij, 94	140	Doxy	Days of productive cough	0.5 (-0.4 to 1.4)
Hueston, 94	23	Erythro	Days of productive cough	-0.4 (-2.4 to 1.6)
King, 96	91	Erythro	Days of sputum production	0.7 (-1.3 to 2.7)

* Positive numbers indicate antibiotics are superior to placebo

Mantel-Haenszel Method (Fixed Effects Model)

	<u>Diseased</u>	<u>Not diseased</u>
Treated (exposed)	a_i	c_i
Not treated (unexposed)	b_i	d_i

$$OR_i = \frac{a_i/c_i}{b_i/d_i} = \frac{a_i \times d_i}{b_i \times c_i}$$

$$InOR_{mh} = \frac{\sum (w_i \times InOR_i)}{\sum w_i}$$

$$\text{variance } InOR_i = \frac{1}{a_i} + \frac{1}{b_i} + \frac{1}{c_i} + \frac{1}{d_i}$$

$$\text{variance } OR_{mh} = \frac{1}{\sum w_i}$$

$$\text{weight}_i = (w_i) = \frac{1}{\text{variance } InOR_i}$$

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Randomized Trials of Antibiotic Rx for acute OM to prevent TM perforation

<u>Study 1</u>	<u>Perforation</u>	<u>No Perforation</u>
Antibiotic	1	114
Placebo	3	116

<u>Study 2</u>	<u>Perforation</u>	<u>No Perforation</u>
Antibiotic	7	65
Placebo	12	65

1. Calculate OR and InOR for each study:

$$OR_1 = \frac{1 \times 116}{3 \times 114} = 0.34 \quad InOR_1 = -1.08$$

$$OR_2 = \frac{7 \times 65}{12 \times 65} = 0.58 \quad InOR_2 = -0.54$$

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Randomized Trials of Antibiotic Rx for acute OM to prevent TM perforation

2. Calculate variance InOR_i for each study:

$$\text{Var } InOR_1 = \frac{1}{1} + \frac{1}{3} + \frac{1}{114} + \frac{1}{116} = 1.35$$

$$\text{Var } InOR_2 = \frac{1}{7} + \frac{1}{12} + \frac{1}{65} + \frac{1}{65} = 0.26$$

3. Calculate w_i for each study:

$$w_1 = \frac{1}{1.35} = 0.74$$

$$w_2 = \frac{1}{0.26} = 3.85$$

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Randomized Trials of Antibiotic Rx
for acute OM to prevent TM perforation

Study 1	Perforation	No Perforation
Antibiotic	1	114
Placebo	3	116

Study 2	Perforation	No Perforation
Antibiotic	7	65
Placebo	12	65

4. Calculate the $w_i \times \ln OR_i$ for each study:

$$w_1 \times \ln OR_1 = 0.74 \times -1.08 = -0.80$$

$$w_2 \times \ln OR_2 = 3.85 \times -0.54 = -2.08$$

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Randomized Trials of Antibiotic Rx
for acute OM to prevent TM perforation

5. Calculate the sum of the w_i

$$w_1 + w_2 = 0.74 + 3.85 = 4.59$$

$$\text{Summary } \ln OR_{mh} = \frac{\sum (w_i \times \ln OR_i)}{\sum w_i} = \frac{-0.80 + -2.08}{4.59} = -0.63$$

$$= OR_{mh} = 0.53$$

$$\text{Calculate variance } OR_{mh} = \frac{1}{\sum w_i} = \frac{1}{4.59} = 0.22$$

8. Calculate 95% CI = $e^{\ln OR_{mh} \pm (1.96 \times \sqrt{\text{variance } \ln OR_{mh}})}$
 $= e^{-0.63 \pm (1.96 \times \sqrt{0.22})} = 0.21 - 1.34$

$$\text{Summary } OR_{mh} = 0.53 \quad (95\% \text{ CI } 0.21 - 1.34)$$

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Dersimonian and Laird Method
(Random Effects Model)

Similar formula to Mantel-Haenszel:

$$\ln OR_{dl} = \frac{\sum (w_i \times \ln OR_i)}{\sum w_i} \quad w_i = \frac{1}{\text{variance}_i + D}$$

- Where D gets larger as the OR (or effect estimate) of the individual studies vary from the summary estimate

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Different types of data

- Different scales (example)
 - Ordinal data
 - Binary data
 - Continuous outcomes

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RCTs in Acute Bronchitis: Different Scales

Study, yr	N	Abx	Outcome	Result
Stott, 76	207	Doxy	Days of Yellow Spit	0.6 (-0.2 to 1.4)
Franks, 84	54	TMP/S	Cough Amount Score	0.2 (-0.2 to 0.6)
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Problem

- How do you combine studies with slightly different outcomes?
- Option 1: - don't do it
- Option 2: Transform all outcomes to an effect size

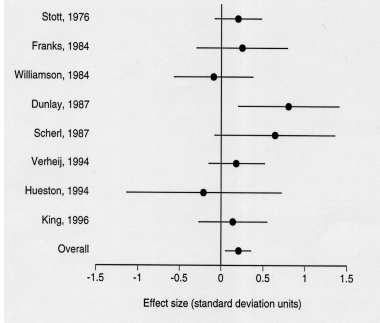
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- Results extracted from study reports may need to be converted to a consistent, or usable, format for analysis .

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Main Result

- Summary ES = 0.21 (95% CI 0.05 to 0.36)

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But...All you need to know is:

- When combined, individual study effect estimates are weighted by their inverse variance
- Variance is related to sample size AND # of events (dichotomous) and precision (continuous)
- Fixed effects just combines all weighted estimates, while random effects "penalizes" estimates for variation between studies

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Thank you
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