

Georgia Salanti and Deborah Caldwell

# Addressing multiple treatments II: multiple-treatments meta-analysis basic methods



Maths Warning!



Πανεπιστήμιο Ιωαννίνων

University of Ioannina



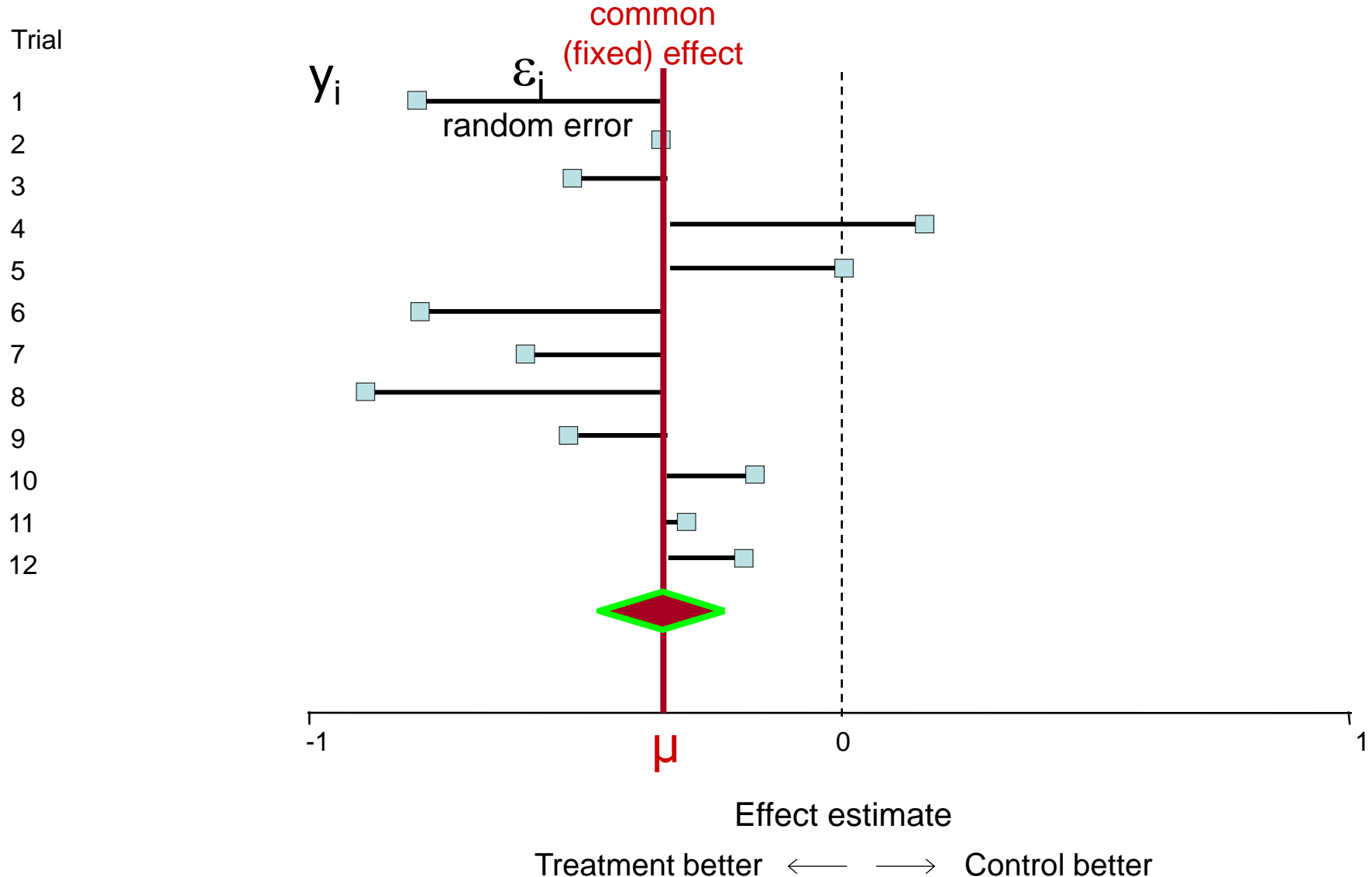
# Why use Bayesian statistics for MTM?

- Bayesian approach is **easier to account for correlations** induced by multi-arm trials
- Estimation of **predictive intervals** is straightforward
- Estimation of **ranking probabilities** is straightforward
- MTM with two-arm trials only  
(or ignoring the correlations)  
Easy with frequentist meta-regression

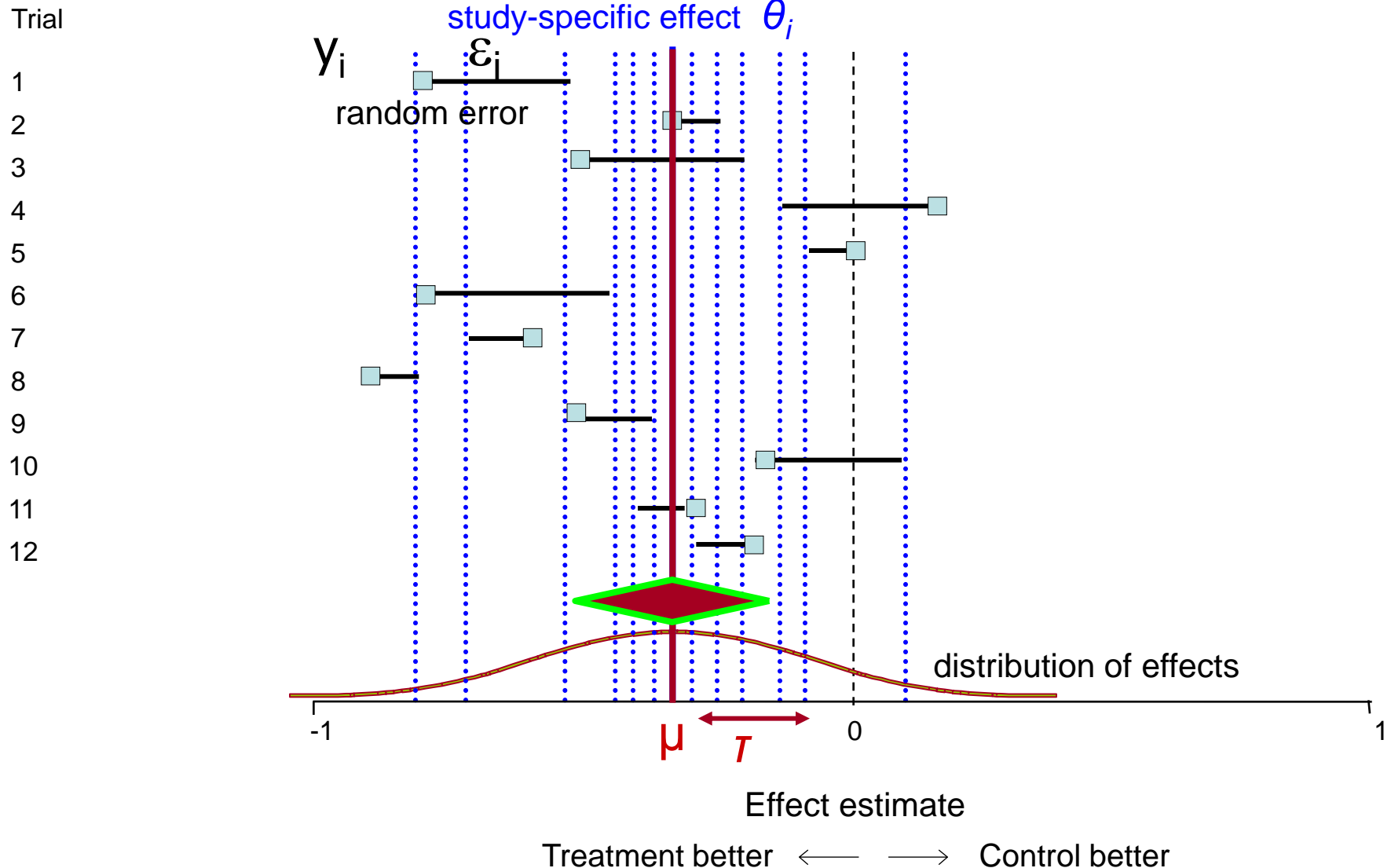
(come to the workshop tomorrow...)



# Fixed effect meta-analysis



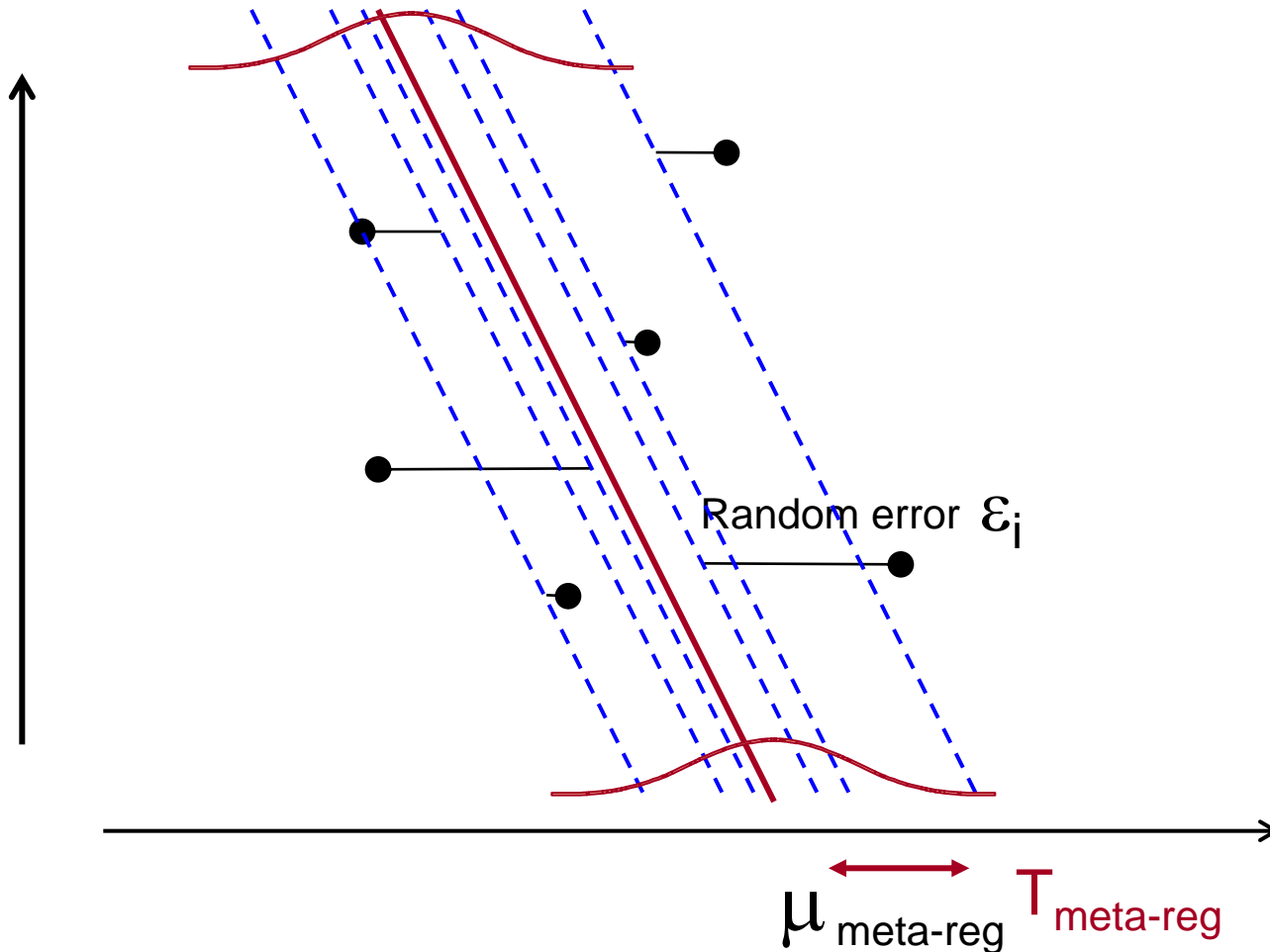
# Random effects meta-analysis



# Random effects meta-regression

$$y_i = \text{intercept} + \text{slope} \times x$$

Explanatory  
variable,  $x$



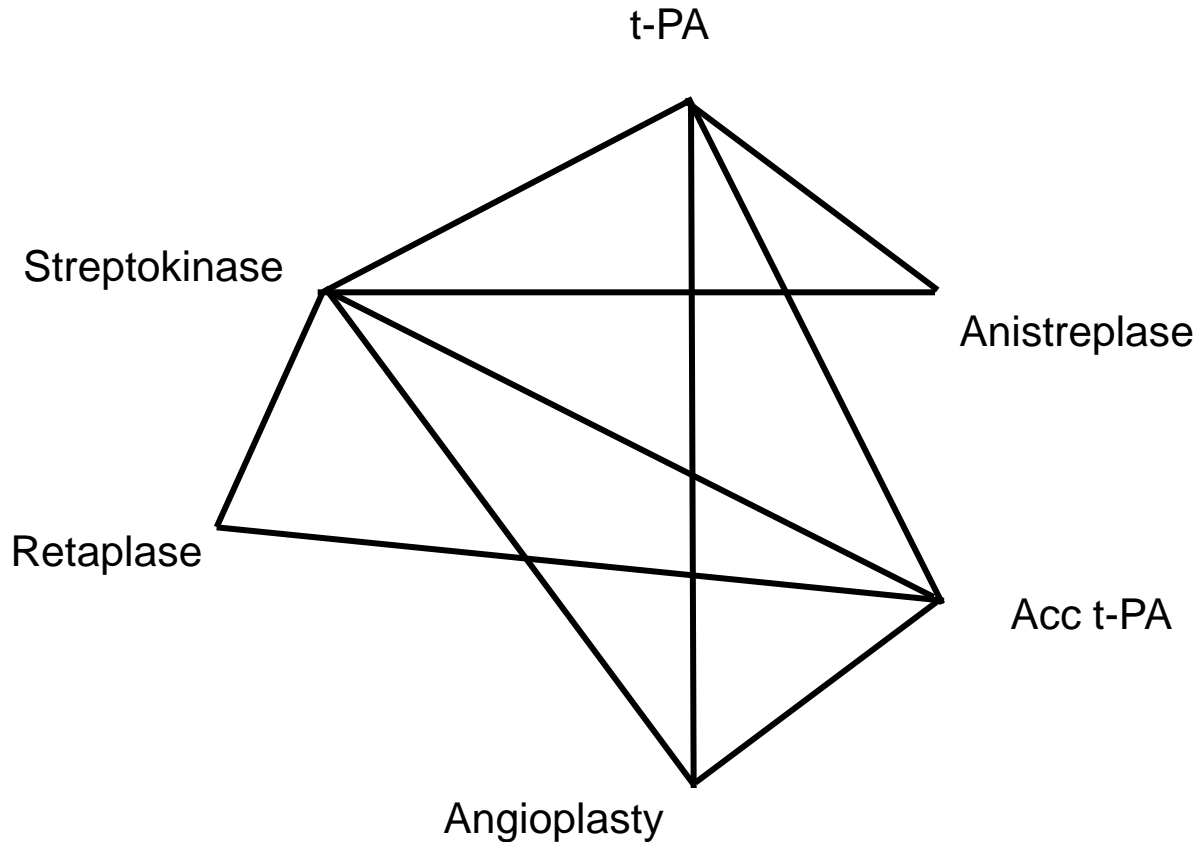
# Meta-regression

- We observe  $y_i$  in each study (e.g. the  $\log(\text{OR})$ )
- Meta-regression using the treatments as ‘covariates’
- AC, AB, BC studies, chose C as *reference*

$$y_i = \mu^{AC} \times (\text{Treat}_i = A) + \mu^{BC} \times (\text{Treat}_i = B)$$

- The AC studies have (1,0), the BC studies (0,1) [*basic*]
- AB studies have (1,-1) [*functional*]
- Please use random effects only

# Parametrisation of the network



Choose basic parameters

Write all other contrasts as linear functions of the basic parameters to build the design matrix

*LOR for death in treatments for MI*

# LOR for death in treatments for MI

$$y_i = \mu^A \text{ t-PA}_i + \mu^B \text{ Anistreplase}_i + \mu^C \text{ Accelerated t-PA}_i + \mu^D \text{ Angioplasty}_i + \mu^E \text{ Reteplase}_i$$

Use as 'covariates'

No. studies	Streptokinase	t-PA	Anistreplase	Acc t-PA	Angioplasty	Reteplase
3	-1	1	0	0	0	0
1	0	0	1	0	0	0
1	0	0	0	1	0	0
3	0	0	0	0	1	0
1	0	0	0	0	0	1
1		-1	1	0	0	0
2		-1	0	0	1	0
2		0	0	-1	1	0
2		0	0	-1	0	1



# LOR for death in treatments for MI

$$y_i = \mu^A \text{ t-PA} + \mu^B \text{ Anistreplase}_i + \mu^C \text{ Accelerated t-PA}_i + \mu^D \text{ Angioplasty}_i + \mu^E \text{ Reteplase}_i$$

$$Y = (\underbrace{\mu^A, \mu^B, \mu^C, \mu^D, \mu^E}_{\text{Vector of LogOR}}) \times \underset{\substack{\uparrow \\ \text{Design} \\ \text{matrix}}}{X} + \underset{\substack{\uparrow \\ \text{Random} \\ \text{effects} \\ \text{matrix}}}{\Delta}$$

Matrix of all observations

$$Y \sim N(\mu X, V)$$

↑  
Variance-covariance  
matrix (for the  
observed LOR)

$$\Delta \sim N(\mathbf{0}, \text{diag}(\tau^2))$$

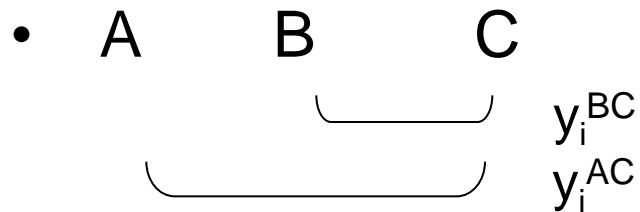
## LOR compared to Streptokinase (RE model)

$$Y = (\mu^A, \mu^B, \mu^C, \mu^D, \mu^E) \times X + \Delta$$

Treatment	LOR(SE)
t-PA	−0.02 (0.03)
Anistreplase	−0.00 (0.03)
Accelerated t-PA	− 0.15 (0.05)
Angioplasty	− 0.43 (0.20)
Reteplase	− 0.11 (0.06)

# What's the problem with multi-arm trials?

- We need to take into account the correlations between the estimates that come from the same study



- The random effects  $(\theta_i^{BC}, \theta_i^{AC})$  that refer to the same trial are correlated as well
- You have to built in *the correlation matrix for the observed effects, **and** the correlation matrix for the random effects*

$$Y \sim N(\mu X, V)$$

$$\Delta \sim N(\mathbf{0}, \text{diag}(\tau^2))$$

# Hypothetical example

Study	No. arms	#	Data	Contrast
i=1	$T_1=2$	1	$y_{1,1}, V_{1,1}$	AB
i=2	$T_2=2$	1	$y_{2,1}, V_{2,1}$	AC
i=3	$T_3=2$	1	$y_{3,1}, V_{3,1}$	BC
i=4	$T_4=3$	2	$y_{4,1}, V_{4,1}$ $y_{4,2}, V_{4,2}$ $\text{cov}(y_{4,1}, y_{4,2})$	AB AC

Basic parameters: AB and AC

Study	No. arms	#	Data	Contrast
i=1	$T_1=2$	1	$y_{1,1}, v_{1,1}$	AB
i=2	$T_2=2$	1	$y_{2,1}, v_{2,1}$	AC
i=3	$T_3=2$	1	$y_{3,1}, v_{3,1}$	BC
i=4	$T_4=3$	2	$y_{4,1}, v_{4,1}$ $y_{4,2}, v_{4,2}$ $\text{cov}(y_{4,1}, y_{4,2})$	AB AC

## Meta-regression

$$\begin{pmatrix} y_{1,1} \\ y_{2,1} \\ y_{3,1} \\ y_{4,1} \\ y_{4,2} \end{pmatrix} = \begin{pmatrix} 1 & 0 \\ 0 & 1 \\ -1 & 1 \\ 1 & 0 \\ 0 & 1 \end{pmatrix} \begin{pmatrix} \mu_{AB} \\ \mu_{AC} \end{pmatrix} + \begin{pmatrix} \beta_{1,1} \\ \beta_{2,1} \\ \beta_{3,1} \\ \beta_{4,1} \\ \beta_{4,2} \end{pmatrix} + \begin{pmatrix} \varepsilon_{1,1} \\ \varepsilon_{2,1} \\ \varepsilon_{3,1} \\ \varepsilon_{4,1} \\ \varepsilon_{4,2} \end{pmatrix}$$

Study	No. arms	#	Data	Contrast
i=1	$T_1=2$	1	$y_{1,1}, v_{1,1}$	AB
i=2	$T_2=2$	1	$y_{2,1}, v_{2,1}$	AC
i=3	$T_3=2$	1	$y_{3,1}, v_{3,1}$	BC
i=4	$T_4=3$	2	$y_{4,1}, v_{4,1}$ $y_{4,2}, v_{4,2}$ $\text{cov}(y_{4,1}, y_{4,2})$	AB AC

$$\begin{pmatrix} y_{1,1} \\ y_{2,1} \\ y_{3,1} \\ y_{4,1} \\ y_{4,2} \end{pmatrix} = \begin{pmatrix} 1 & 0 \\ 0 & 1 \\ -1 & 1 \\ 1 & 0 \\ 0 & 1 \end{pmatrix} \begin{pmatrix} \mu_{AB} \\ \mu_{AC} \end{pmatrix} + \begin{pmatrix} \beta_{1,1} \\ \beta_{2,1} \\ \beta_{3,1} \\ \beta_{4,1} \\ \beta_{4,2} \end{pmatrix} + \begin{pmatrix} \varepsilon_{1,1} \\ \varepsilon_{2,1} \\ \varepsilon_{3,1} \\ \varepsilon_{4,1} \\ \varepsilon_{4,2} \end{pmatrix}$$

Take into account correlation  
in observations

$$\begin{pmatrix} \varepsilon_{1,1} \\ \varepsilon_{2,1} \\ \varepsilon_{3,1} \\ \varepsilon_{4,1} \\ \varepsilon_{4,2} \end{pmatrix} \sim N \left( \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \end{pmatrix}, \begin{pmatrix} v_{1,1} & 0 & 0 & 0 & 0 \\ 0 & v_{2,1} & 0 & 0 & 0 \\ 0 & 0 & v_{3,1} & 0 & 0 \\ 0 & 0 & 0 & v_{4,1} & \text{cov}(y_{4,1}, y_{4,2}) \\ 0 & 0 & 0 & \text{cov}(y_{4,1}, y_{4,2}) & v_{4,2} \end{pmatrix} \right)$$

Study	No. arms	#	Data	Contrast
i=1	$T_1=2$	1	$y_{1,1}, v_{1,1}$	AB
i=2	$T_2=2$	1	$y_{2,1}, v_{2,1}$	AC
i=3	$T_3=2$	1	$y_{3,1}, v_{3,1}$	BC
i=4	$T_4=3$	2	$y_{4,1}, v_{4,1}$ $y_{4,2}, v_{4,2}$ $\text{cov}(y_{4,1}, y_{4,2})$	AB AC

$$\begin{pmatrix} y_{1,1} \\ y_{2,1} \\ y_{3,1} \\ y_{4,1} \\ y_{4,2} \end{pmatrix} = \begin{pmatrix} 1 & 0 \\ 0 & 1 \\ -1 & 1 \\ 1 & 0 \\ 0 & 1 \end{pmatrix} \begin{pmatrix} \mu_{AB} \\ \mu_{AC} \end{pmatrix} + \boxed{\begin{pmatrix} \beta_{1,1} \\ \beta_{2,1} \\ \beta_{3,1} \\ \beta_{4,1} \\ \beta_{4,2} \end{pmatrix}} + \begin{pmatrix} \varepsilon_{1,1} \\ \varepsilon_{2,1} \\ \varepsilon_{3,1} \\ \varepsilon_{4,1} \\ \varepsilon_{4,2} \end{pmatrix}$$

Take into account correlation  
in random effects

$$\begin{pmatrix} \beta_{1,1} \\ \beta_{2,1} \\ \beta_{3,1} \\ \beta_{4,1} \\ \beta_{4,2} \end{pmatrix} \sim N \left( \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \end{pmatrix}, \begin{pmatrix} \tau_{AB}^2 & 0 & 0 & 0 & 0 \\ 0 & \tau_{AC}^2 & 0 & 0 & 0 \\ 0 & 0 & \tau_{BC}^2 & 0 & 0 \\ 0 & 0 & 0 & \tau_{AB}^2 & \text{cov}(\beta_{4,1}, \beta_{4,2}) \\ 0 & 0 & 0 & \text{cov}(\beta_{4,1}, \beta_{4,2}) & \tau_{AC}^2 \end{pmatrix} \right)$$

# How to fit such a model?

- MLwiN
- SAS, R
- STATA using metan

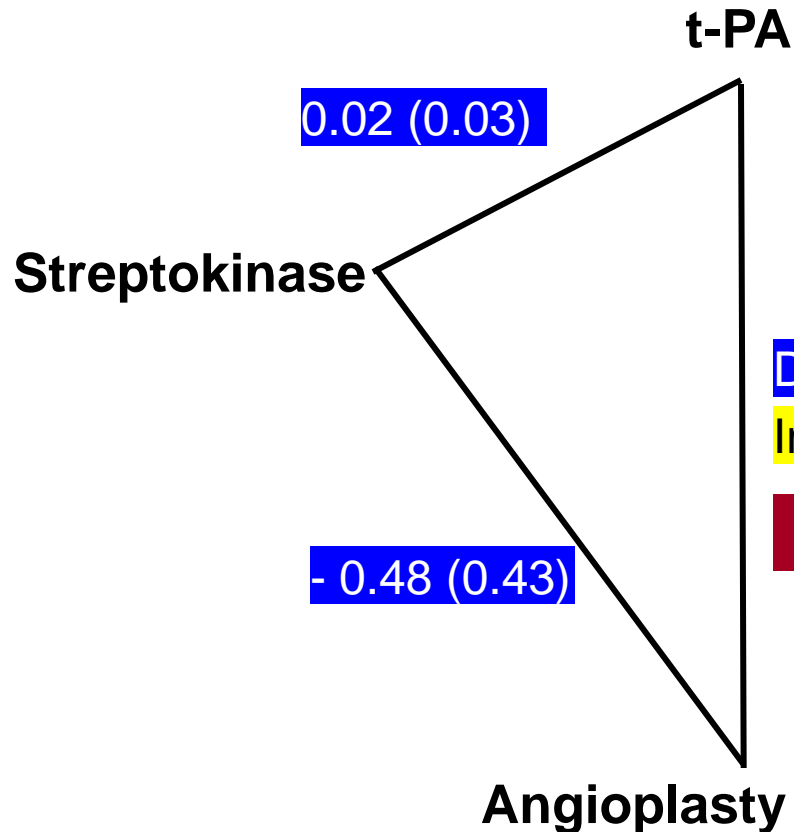
*Review of statistical methodology in Salanti G, Higgins JP, Ades AE, Ioannidis JP. 2008. Evaluation of networks of randomized trials. Statistical Methods in Medical Research 17:279-301.*



# Inconsistency

*LOR (SE) for MI*

Calculate a difference  
between direct and  
indirect estimates



Direct t-PA vs Angioplasty = - 0.41 (0.36)

Indirect t-PA vs Angioplasty = - 0.46 (0.18)

Inconsistency Factor IF = 0.05

# Inconsistency - Heterogeneity

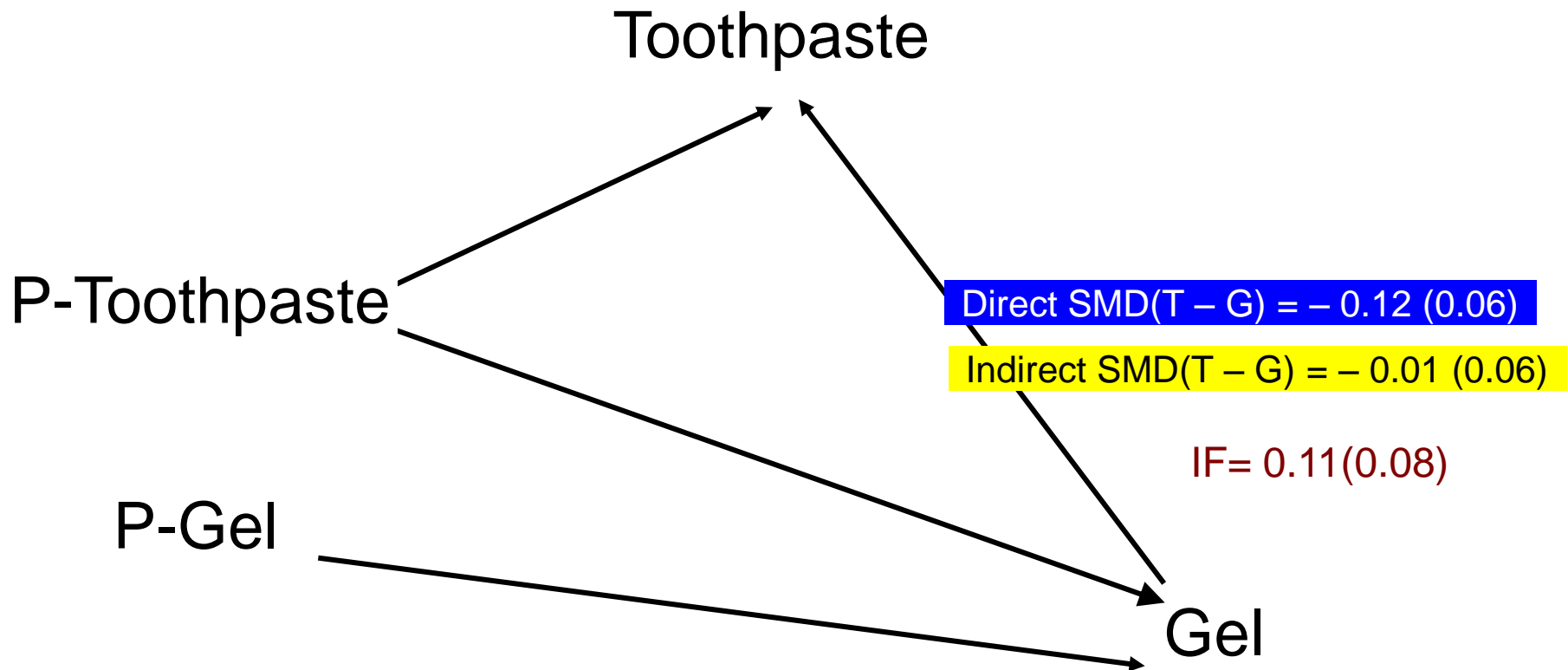
- *Heterogeneity*: 'excessive' discrepancy among study-specific effects
- *Inconsistency*: it is the excessive discrepancy among source-specific effects (direct and indirect)
- In 3 cases out of 44 there was an important discrepancy between direct/indirect effect.

*Glenny et al HTA 2005*

# What can cause inconsistency?

*Inappropriate common comparator*

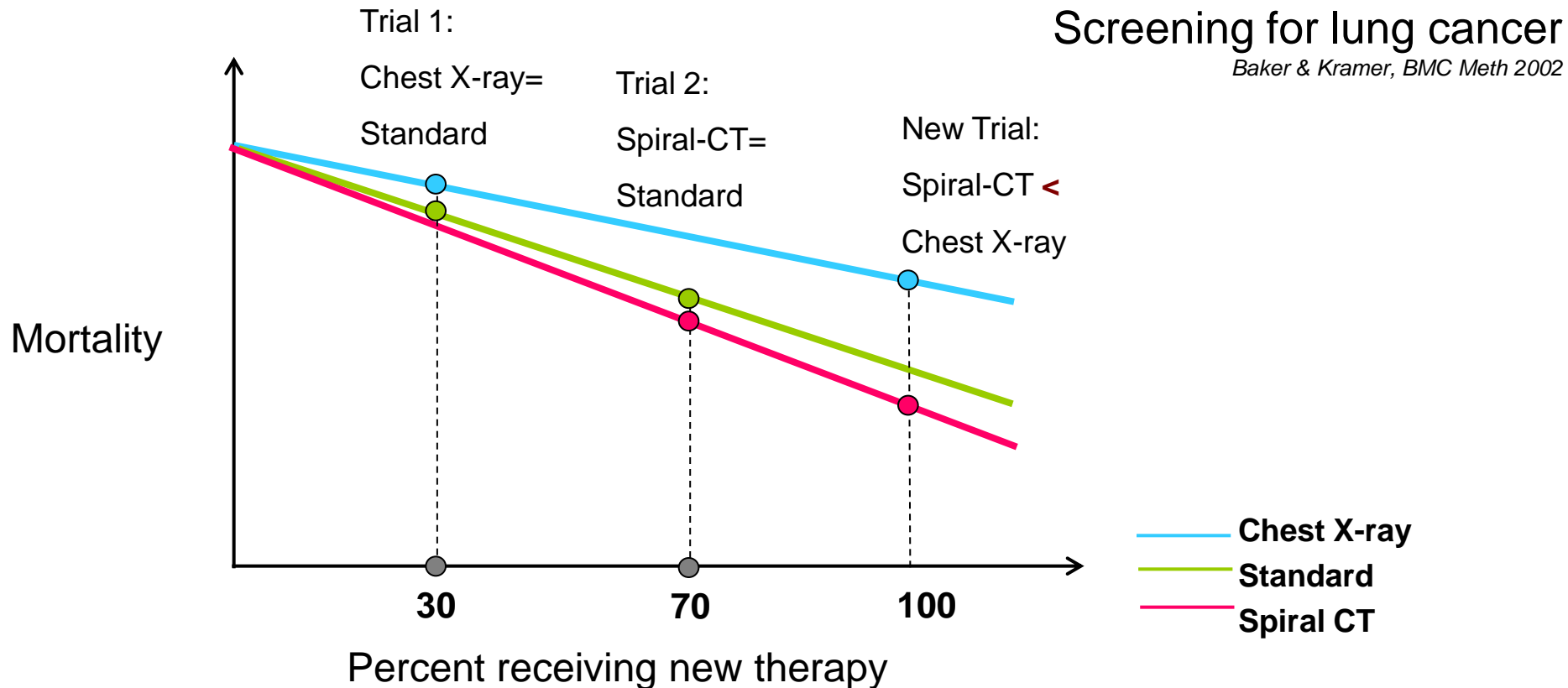
Compare Fluoride treatments in preventing dental caries



I cannot learn about Toothpaste versus Gel through Placebo!

# What can cause inconsistency?

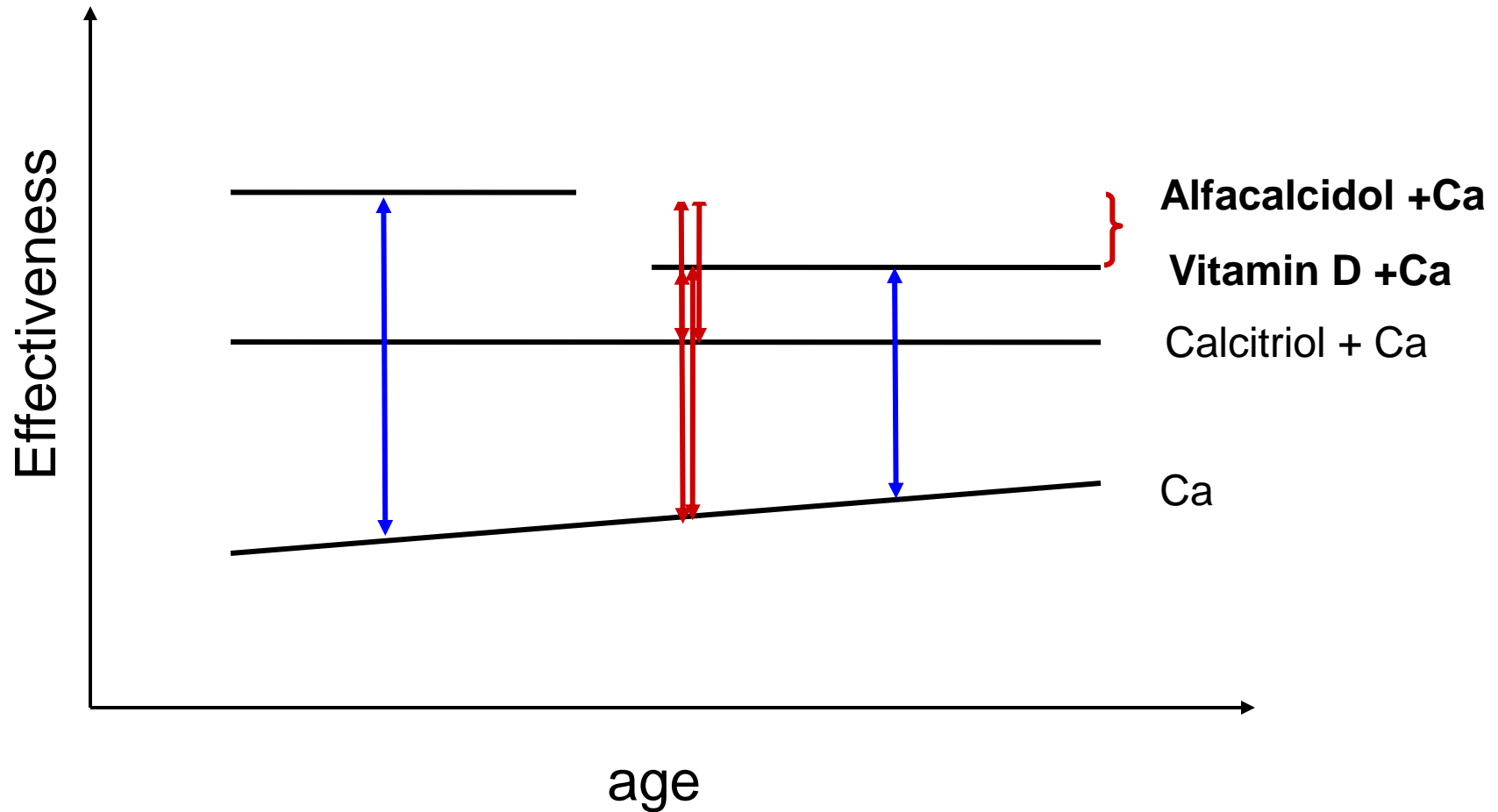
*Confounding by trial characteristics*



A new therapy (possibly unreported in the trials) decreases the mortality but in **different rates** for the three screening methods

# What can cause inconsistency?

*Confounding by trial characteristics*



Different characteristics across comparisons may cause inconsistency

# Assumptions of MTM

- There is **not confounding** by trial characteristics that are related to both the comparison being made and the magnitude of treatment difference
- The trials in two different comparisons are **exchangeable** (other than interventions being compared)
- Equivalent to the assumption '***the unobserved treatment is missing at random***'
  - *Is this plausible?*
  - *Selection of the comparator is not often random!*

# Inconsistency

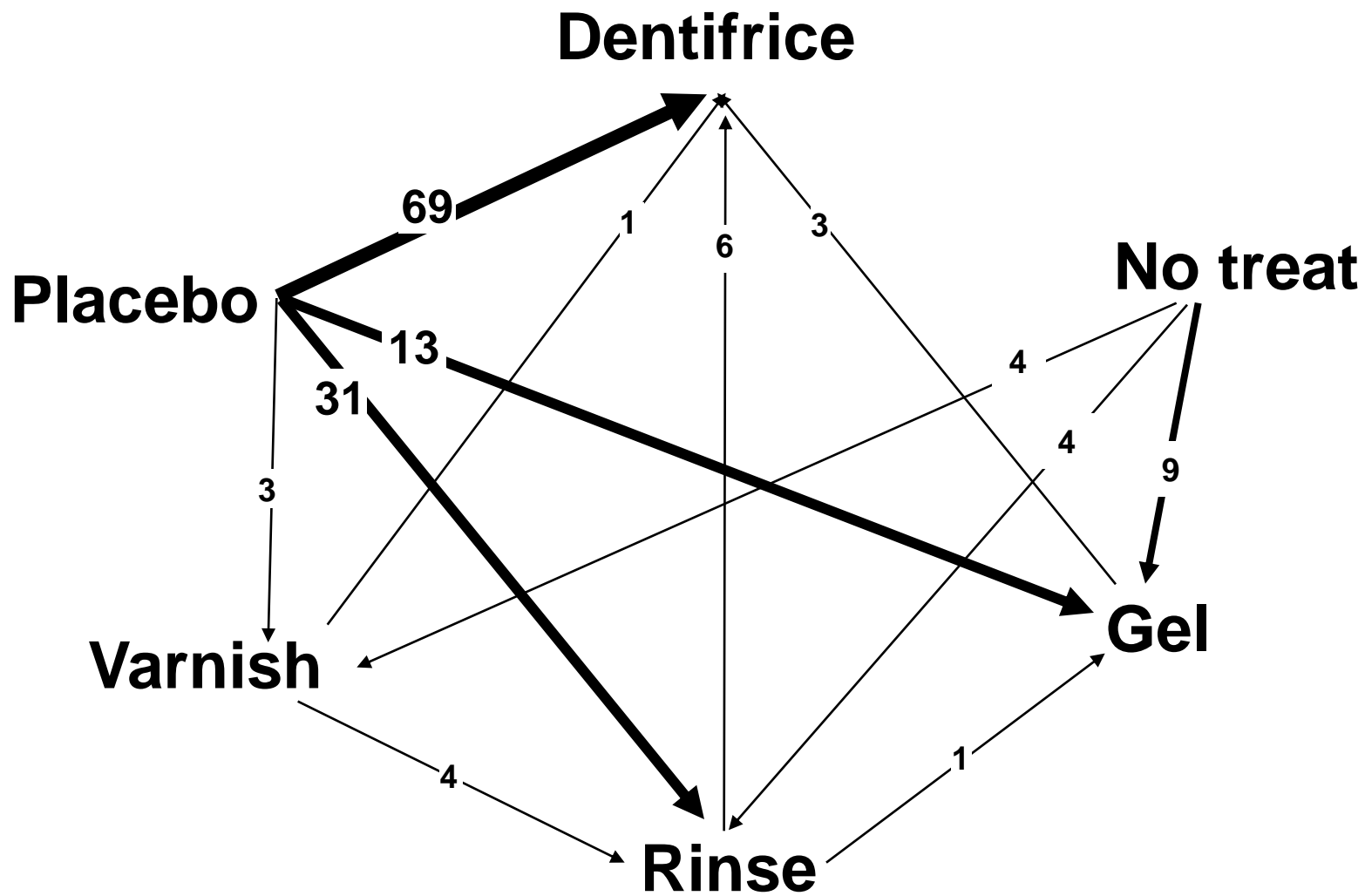
## *Detecting*

- Check the distribution of important characteristics per treatment comparison
  - Usually unobserved....
  - **Time** (of randomization, of recruitment) might be associated with changes to the background risk that may violate the assumptions of MTM
- Get a taste by looking for inconsistency in closed loops
- Fit a model that relaxes consistency
  - Add an extra '*random effect*' per loop (Lu & Ades JASA 2005)

# Compare the characteristics!

No. studies	T	G	R	V	P	Fup	Baseline	Year	Water F (yes/no)
69						2.6	11.8	1968	0.2
13						2.3	3.8	1973	0.2
30						2.4	5.9	1973	0.1
3						2.3	2.7	1983	0
3						2.7	NA	1968	0.66
6						2.8	14.7	1969	0
1						2	0.9	1978	0
1						1	NA	1977	0
1						3	7.4	1991	NA
4						2.5	7.6	1981	0.33

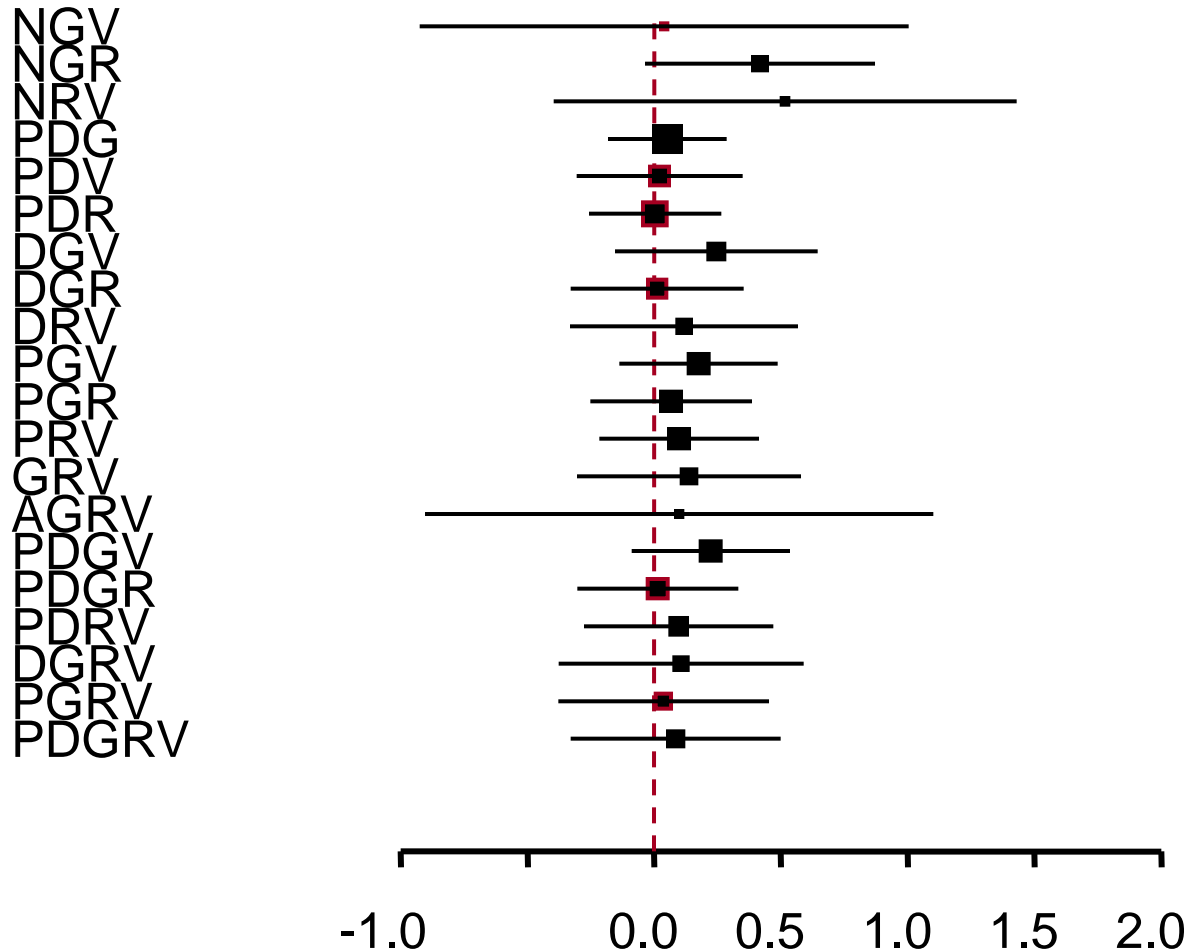




# Evaluation of concordance within closed loops

Estimates with 95% confidence intervals

Closed loops



R routine in <http://www.dhe.med.uoi.gr/software.htm>

Salanti G, Marinho V, Higgins JP: **A case study of multiple-treatments meta-analysis demonstrates that covariates should be considered.** *J Clin Epidemiol* 2009, **62**: 857-864.

# More assumptions of MTM!

- *Appropriate modelling of data* (sampling distributions)
- *Normality* of true effects in a random-effects analysis
- *Comparability of studies*
  - exchangeability in all aspects other than particular treatment comparison being made
- *Equal heterogeneity variance in each comparison*
  - not strictly necessary

# References

- Caldwell DM, Ades AE, Higgins JP. 2005. Simultaneous comparison of multiple treatments: combining direct and indirect evidence. *BMJ* 331:897-900.
- Caldwell DM, Gibb DM, Ades AE. 2007. Validity of indirect comparisons in meta-analysis. *Lancet* 369:270.
- Caldwell DM, Welton NJ, Ades AE. 2010. Mixed treatment comparison analysis provides internally coherent treatment effect estimates based on overviews of reviews and can reveal inconsistency. *J Clin Epidemiol*.
- Dias S, Welton N, Marinho V, Salanti G, Ades A. 2010. Estimation and adjustment of Bias in randomised evidence using Mixed Treatment Comparison Meta-analysis. *Journal of the Royal Statistical Society (A)* 173.
- Glenny AM, Altman DG, Song F, Sakarovitch C, Deeks JJ, D'Amico R, Bradburn M, Eastwood AJ. 2005. Indirect comparisons of competing interventions. *Health Technol Assess* 9:1-iv
- Salanti G, Higgins JP, Ades AE, Ioannidis JP. 2008. Evaluation of networks of randomized trials. *Statistical Methods in Medical Research* 17:279-301.
- Salanti G, Marinho V, Higgins JP. 2009. A case study of multiple-treatments meta-analysis demonstrates that covariates should be considered. *Journal of Clinical Epidemiology* 62:857-864.
- Song F, Altman DG, Glenny AM, Deeks JJ. 2003. Validity of indirect comparison for estimating efficacy of competing interventions: empirical evidence from published meta-analyses. *BMJ* 326:472.
- Song F, Harvey I, Lilford R. 2008. Adjusted indirect comparison may be less biased than direct comparison for evaluating new pharmaceutical interventions. *Journal of Clinical Epidemiology* 61:455-463.
- Sutton A, Ades AE, Cooper N, Abrams K. 2008. Use of indirect and mixed treatment comparisons for technology assessment. *Pharmacoeconomics* 26:753-767.