A comparison of one vs two stage dose-response meta-analysis

Alessio Crippa^{1,2}, Nicola Orsini^{1,2} ¹Unit of Nutritional Epidemiology, Institute of Environmental Medicine, Karolinska Institutet, Stockholm, Sweden ²Unit of Biostatistics, Institute of Environmental Medicine, Karolinska Institutet, Stockholm, Sweden

Conclusions

One and two stage dose-response meta-analysis provide the same curves.

A one stage approach, however, allows:

- simultaneous estimation of both study-specific and pooled trend;
- inclusion of additional studies with only few data points;
- estimation of more flexible curves;
- interaction analysis with study level covariates;
- comparison of alternative models using information criteria.



Figure 1. Predicted dose-response curves for a dose-response metaanalysis between coffee consumption and all-cause mortality. Models compared are restricted cubic splines with 3 knots, exclusion of studies with less than three exposure categories, and 4 knots.

Introduction

Dose-response meta-analyses are increasingly popular. A standard approach is a two-stage analysis: studyspecific trend estimation and pooling of study-specific dose-response coefficients. An alternative one stage technique has never been fully described.

Aims

To investigate analogies and dissimilarities between the one- and two-stage dose-response meta-analysis using a real example.

One vs two stage approaches

We use restricted cubic splines (RCSpline) with k = 3knots to examine the shape of the association between coffee consumption and mortality, based on I studies (i =1, ..., I)

Two-stage: $ln(RR_{ij}) = \beta_{1i}x_{1ij} + \beta_{2i}x_{2ij} + \varepsilon_{ij}, j = 1, ..., n_i$ $(\hat{\beta}_{1i}, \hat{\beta}_{2i})^T \sim N\left((\bar{\beta}_1, \bar{\beta}_2)^T, \Psi\right)$

One-stage: $ln(RR_{ij}) = (\bar{\beta}_1 + u_{1i})x_1$ $\varepsilon_{ij} \sim N(0, \Sigma_i); \quad (u_{1i}, u_{2i})^T \sim N(0, \Psi)$

In the two-stage approach, studies with less than three exposure categories are excluded (e.g. second study in Table 1).



Karolinska Institutet Alessio Crippa Institute of Environmental Medicine Nobelsväg 13 P.O. Box 210 SE-171 77, Stockholm

$$x_{1ij} + (\bar{\beta}_2 + u_{2i})x_{2ij} + \varepsilon_{ij}$$

id	author	cases	n	dose	RR	(95% CI)
1	LeGrady et al.	57	249	0.5	1.00	-
		136	655	2.5	0.75	(0.57, 0.99)
		144	619	4.5	0.84	(0.64-1.10)
		115	387	6.5	1.09	(0.83-1.43)
2	Nilsson et al.			0.0	1.00	-
				2.0	0.99	(0.86-1.14)

Table 1. Aggregated data for two studies included in a dose-response meta-analysis between coffee consumption and all-cause mortality.

Results

Excluding two studies with less than three exposure categories, both approaches provide the same coefficients estimates (Table 2).

The one-stage analysis avoids exclusion of such studies (Table 2) and may evaluate even more flexible curves, e.g. more knots in a spline model. The RCSpline with k=4 knots, however, has a higher AIC, suggesting that the simpler (k=3) properly fits the data (Figure 1). The same comparison can not be performed in a two-stage analysis.

analysis	excl.	$\widehat{\beta}_1$	$\widehat{\beta}_2$	$Var(\widehat{\overline{\beta}}_1)$	$Cov(\widehat{\overline{\beta}}_1\widehat{\overline{\beta}}_2)$	$Var(\widehat{\overline{\beta}}_2)$
two-stage	yes	-0.09216	0.08666	0.00019	-0.00018	0.00019
one-stage	yes	-0.09217	0.08665	0.00019	-0.00018	0.00019
one-stage	no	-0.08357	0.07714	0.00020	-0.00018	0.00017

Table 2. Dose-response coefficients estimates for one- and two-stage dose-response meta-analysis between coffee consumption and allcause mortality.



