#### Design of dose-escalation trials

R. A. Bailey



r.a.bailey@qmul.ac.uk

Belgian Statistical Society Antwerp October 2007

#### Abstract

In one form of dose-escalation trial, several cohorts of subjects are recruited. Each cohort takes part at a different time period. The doses are ordinally labelled  $0, 1, \ldots$  Because higher doses may have more adverse side-effects, no subject can be exposed to dose i until some information is obtained about the effect of dose i-1.

One possibility is to use dose i for everyone in cohort i. Then there is no blinding; moreover, dose effects are completely confounded with cohort effects and period effects.

A modification of this uses a certain number of placebo subjects in each cohort. If there are no cohort effects then the proportion of placebo in each cohort should be such that the design is equireplicate if it proceeds to the planned largest dose. If there are cohort effects, then more precise comparisons between doses can be made if half of each cohort receives placebo.

I shall discuss a new design that does at least as well as both of these, whether or not there is a cohort effect.

## Design of the TeGenero trial

Cohort	TGN141	Placebo	
	Dose	Number of	Number of
	mg/kg bodyweight	Subjects	Subjects
1	0.1	6	2
2	0.5	6	2
3	2.0	6	2
4	5.0	6	2

## What happened to Cohort 1 on 13 March 2006

Healthy	Randomised	Time of	Time of
Volunteer	to	intravenous	transfer to
		administration	critical care
A	TGN1412 8.4mg	0800	2400
В	Placebo	0810	
С	TGN1412 6.8mg	0820	2350
D	TGN1412 8.8mg	0830	0030
Е	TGN1412 8.2mg	0840	2040
F	TGN1412 7.2mg	0850	0050
G	TGN1412 8.2mg	0900	0100
Н	Placebo	0910	

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

Recommendations include

generic issues

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects
   (sudden adverse effects → do not dose further subjects;
   delayed adverse effects → ill subjects can be treated one by one)

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects (sudden adverse effects → do not dose further subjects; delayed adverse effects → ill subjects can be treated one by one)
- preclinical / clinical interface

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects (sudden adverse effects → do not dose further subjects; delayed adverse effects → ill subjects can be treated one by one)
- preclinical / clinical interface
- protocol

Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects (sudden adverse effects → do not dose further subjects; delayed adverse effects → ill subjects can be treated one by one)
- preclinical / clinical interface
- protocol
- sequential choice of dose



Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects
   (sudden adverse effects → do not dose further subjects;
   delayed adverse effects → ill subjects can be treated one by one)
- preclinical / clinical interface
- protocol
- sequential choice of dose
- allocation of ordinal doses to cohorts.



Published in March 2007 on http://www.rss.org.uk/main.asp?page=2816

- generic issues
- ► risk (quantification; novel type of medicine; public debate)
- sharing information on adverse events (usable database)
- ▶ proper interval between dosing subjects (sudden adverse effects → do not dose further subjects; delayed adverse effects → ill subjects can be treated one by one)
- preclinical / clinical interface
- protocol
- sequential choice of dose
- allocation of ordinal doses to cohorts.



Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

If all responses are uncorrelated with variance  $\sigma^2$  then Variance (dose i- placebo) in cohort i is  $\left(\frac{1}{6}+\frac{1}{2}\right)\sigma^2=\frac{2}{3}\sigma^2$ 

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

If all responses are uncorrelated with variance  $\sigma^2$  then Variance (dose i- placebo) in cohort i is  $\left(\frac{1}{6}+\frac{1}{2}\right)\sigma^2=\frac{2}{3}\sigma^2$ 

From the protocol: "data of subjects having received placebo will be pooled in one group for analyses"

Cohort	TGN1412		Placebo
	Dose Number		Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

If all responses are uncorrelated with variance  $\sigma^2$  then Variance (dose i- placebo) in cohort i is  $\left(\frac{1}{6}+\frac{1}{2}\right)\sigma^2=\frac{2}{3}\sigma^2$ 

From the protocol: "data of subjects having received placebo will be pooled in one group for analyses"

Variance (dose 
$$i-$$
 placebo) is  $\left(\frac{1}{6} + \frac{1}{8}\right)\sigma^2 = \frac{7}{24}\sigma^2$ 

Cohort	TGN1412		Placebo
	Dose Number		Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

If all responses are uncorrelated with variance  $\sigma^2$  then Variance (dose i- placebo) in cohort i is  $\left(\frac{1}{6}+\frac{1}{2}\right)\sigma^2=\frac{2}{3}\sigma^2$ 

From the protocol: "data of subjects having received placebo will be pooled in one group for analyses"

Variance (dose i- placebo) is  $\left(\frac{1}{6} + \frac{1}{8}\right) \sigma^2 = \frac{7}{24} \sigma^2$  if there are no cohort effects

Cohort	TGN1412		Placebo
	Dose Number		Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

If all responses are uncorrelated with variance  $\sigma^2$  then Variance (dose i- placebo) in cohort i is  $\left(\frac{1}{6}+\frac{1}{2}\right)\sigma^2=\frac{2}{3}\sigma^2$ 

From the protocol: "data of subjects having received placebo will be pooled in one group for analyses"

Variance (dose i – placebo) is  $\left(\frac{1}{6} + \frac{1}{8}\right) \sigma^2 = \frac{7}{24} \sigma^2$  if there are no cohort effects

Variance (dose i- dose j) is  $\left(\frac{1}{6}+\frac{1}{6}\right)\sigma^2=\frac{1}{3}\sigma^2$  if there are no cohort effects

▶ Different types of people can volunteer at different times.

- ▶ Different types of people can volunteer at different times.
- ► There may be changes in the ambient conditions, eg temperature, pollutants, pollens.

- ▶ Different types of people can volunteer at different times.
- ► There may be changes in the ambient conditions, eg temperature, pollutants, pollens.
- ► The staff running the trial, or analysing the samples, may change.

- ▶ Different types of people can volunteer at different times.
- ► There may be changes in the ambient conditions, eg temperature, pollutants, pollens.
- ► The staff running the trial, or analysing the samples, may change.

- ▶ Different types of people can volunteer at different times.
- ► There may be changes in the ambient conditions, eg temperature, pollutants, pollens.
- ► The staff running the trial, or analysing the samples, may change.

There have been many trials, in many topics, where, with hindsight, cohort effects swamp treatment effects.

#### Analysis of the TeGenero trial with cohort effects

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{6} + \frac{1}{2}\right)\sigma^2 = \frac{2}{3}\sigma^2$ .

#### Analysis of the TeGenero trial with cohort effects

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{6} + \frac{1}{2}\right)\sigma^2 = \frac{2}{3}\sigma^2$ .

Estimator of (dose 
$$i$$
 – dose  $j$ ) = [estimator of (dose  $i$  – placebo) in cohort  $i$ ] – [estimator of (dose  $j$  – placebo) in cohort  $j$ ]

#### Analysis of the TeGenero trial with cohort effects

Cohort	TGN1412		Placebo
	Dose Number		Number
1	1	6	2
2	2	6	2
3	3	6	2
4	4	6	2

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{6} + \frac{1}{2}\right)\sigma^2 = \frac{2}{3}\sigma^2$ .

Estimator of (dose 
$$i$$
 – dose  $j$ ) = [estimator of (dose  $i$  – placebo) in cohort  $i$ ] – [estimator of (dose  $j$  – placebo) in cohort  $j$ ]

So variance (dose 
$$i - \text{dose } j$$
) =  $\left(\frac{2}{3} + \frac{2}{3}\right) \sigma^2 = \frac{4}{3} \sigma^2$ .

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{4} + \frac{1}{4}\right)\sigma^2 = \frac{1}{2}\sigma^2$ 

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{4} + \frac{1}{4}\right)\sigma^2 = \frac{1}{2}\sigma^2 < \frac{2}{3}\sigma^2$ .

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{4} + \frac{1}{4}\right)\sigma^2 = \frac{1}{2}\sigma^2 < \frac{2}{3}\sigma^2$ .

So variance (dose 
$$i$$
 – dose  $j$ ) =  $\left(\frac{1}{2} + \frac{1}{2}\right) \sigma^2 = \sigma^2$ 

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{4} + \frac{1}{4}\right)\sigma^2 = \frac{1}{2}\sigma^2 < \frac{2}{3}\sigma^2$ .

So variance (dose 
$$i$$
 – dose  $j$ ) =  $\left(\frac{1}{2} + \frac{1}{2}\right)\sigma^2 = \sigma^2 < \frac{4}{3}\sigma^2$ .

Cohort	TGN1412		Placebo
	Dose	Number	Number
1	1	4	4
2	2	4	4
3	3	4	4
4	4	4	4

Variance (dose 
$$i$$
 – placebo) in cohort  $i = \left(\frac{1}{4} + \frac{1}{4}\right)\sigma^2 = \frac{1}{2}\sigma^2 < \frac{2}{3}\sigma^2$ .

So variance (dose 
$$i$$
 – dose  $j$ ) =  $\left(\frac{1}{2} + \frac{1}{2}\right)\sigma^2 = \sigma^2 < \frac{4}{3}\sigma^2$ .

The TeGenero design is inadmissible because everything can be estimated, from the same resources, with smaller variance, by another design.

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n cohorts of m subjects each.

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n cohorts of m subjects each.

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n cohorts of m subjects each.

Cohort 1 subjects may receive only dose 1 or placebo.

In Cohort i, some subjects receive dose i; no subject receives dose j if j > i.

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n cohorts of m subjects each.

Cohort 1 subjects may receive only dose 1 or placebo.

In Cohort i, some subjects receive dose i; no subject receives dose j if j > i.

Put  $s_{ik}$  = number of subjects who get dose i in period k. Then

$$s_{ik} > 0$$
 if  $i = k$   
 $s_{ik} = 0$  if  $i > k$ 

Assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i + \beta_k$ , and that responses are uncorrelated with common variance  $\sigma^2$ .

Assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i + \beta_k$ , and that responses are uncorrelated with common variance  $\sigma^2$ .

"Variance (dose i - dose j)" means  $\text{Var}(\widehat{\tau}_i - \widehat{\tau}_j)$ .

Assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i + \beta_k$ , and that responses are uncorrelated with common variance  $\sigma^2$ .

"Variance (dose i - dose j)" means  $\text{Var}(\widehat{\tau}_i - \widehat{\tau}_j)$ .

Assess designs by looking at the pairwise variances.

Assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i + \beta_k$ , and that responses are uncorrelated with common variance  $\sigma^2$ .

"Variance (dose i - dose j)" means  $\text{Var}(\widehat{\tau}_i - \widehat{\tau}_j)$ .

Assess designs by looking at the pairwise variances.

Doubling the number of subjects — halving all variances,

Assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i + \beta_k$ , and that responses are uncorrelated with common variance  $\sigma^2$ .

"Variance (dose i - dose j)" means  $\text{Var}(\widehat{\tau}_i - \widehat{\tau}_j)$ .

Assess designs by looking at the pairwise variances.

Doubling the number of subjects → halving all variances,

so define the scaled variance  $v_{ij}$  to be

$$\frac{\text{Variance } (\text{dose } i - \text{ dose } j) \times \text{number of subjects}}{\sigma^2}$$

- ightharpoonup only doses 0 and k in period k
- equal replication overall.

- only doses 0 and k in period k
- equal replication overall.

$$s_{ik} = \begin{cases} \frac{m}{n+1} & \text{if } i = 0\\ \frac{nm}{n+1} & \text{if } 0 < i = k\\ 0 & \text{otherwise.} \end{cases}$$

- only doses 0 and k in period k
- equal replication overall.

$$s_{ik} = \begin{cases} \frac{m}{n+1} & \text{if } i = 0 \\ \frac{nm}{n+1} & \text{if } 0 < i = k \end{cases}$$
 Example:  $n = 3, m = 8$ 

$$\frac{\text{Dose}}{\text{Cohort 1}} \begin{vmatrix} 0 & 1 & 2 & 3 \\ 2 & 6 & 0 & 0 \\ \text{Cohort 2} \begin{vmatrix} 2 & 6 & 0 & 0 \\ 2 & 0 & 6 & 0 \\ \text{Cohort 3} \end{vmatrix} = 0$$

- only doses 0 and k in period k
- equal replication overall.

$$s_{ik} = \begin{cases} \frac{m}{n+1} & \text{if } i = 0 \\ \frac{nm}{n+1} & \text{if } 0 < i = k \end{cases}$$
 Example:  $n = 3, m = 8$ 

$$\frac{\text{Dose}}{\text{Cohort 1}} = \frac{0.1 + 0.2 + 0.3}{0.2 + 0.2} = \frac{0.1 + 0.2 + 0.2}{0.2 + 0.2} = \frac{0.1 + 0.2 + 0.2}{0.2} = \frac{0.1 + 0.2}{0.2} = \frac{0.2 + 0.2}$$

$$v_{0i} = (n+1)^2$$
  
 $v_{ij} = 2(n+1)^2$ 

- ightharpoonup only doses 0 and k in period k
- ▶ minimize pairwise variances if there are period effects.

- only doses 0 and k in period k
- minimize pairwise variances if there are period effects.

$$s_{ik} = \begin{cases} \frac{m}{2} & \text{if } i = 0\\ \frac{m}{2} & \text{if } 0 < i = k\\ 0 & \text{otherwise.} \end{cases}$$

- only doses 0 and k in period k
- ▶ minimize pairwise variances if there are period effects.

$$s_{ik} = \begin{cases} \frac{m}{2} & \text{if } i = 0 \\ \frac{m}{2} & \text{if } 0 < i = k \end{cases}$$
 Example:  $n = 3, m = 8$ 

$$\frac{\text{Dose}}{\text{Cohort 1}} \begin{vmatrix} 0 & 1 & 2 & 3 \\ 4 & 4 & 0 & 0 \\ \text{Cohort 2} \end{vmatrix} \begin{vmatrix} 0 & 1 & 2 & 3 \\ 4 & 0 & 4 & 0 \\ \text{Cohort 3} \end{vmatrix}$$

- only doses 0 and k in period k
- minimize pairwise variances if there are period effects.

$$s_{ik} = \begin{cases} \frac{m}{2} & \text{if } i = 0 \\ \frac{m}{2} & \text{if } 0 < i = k \end{cases}$$
 Example:  $n = 3, m = 8$ 

$$\frac{\text{Dose}}{\text{Cohort 1}} \begin{vmatrix} 0 & 1 & 2 & 3 \\ 4 & 4 & 0 & 0 \\ \text{Cohort 2} \end{vmatrix} \begin{vmatrix} 0 & 4 & 0 \\ 4 & 0 & 4 \end{vmatrix}$$

$$\frac{\text{Cohort 3}}{\text{Cohort 3}} \begin{vmatrix} 0 & 4 & 0 \\ 4 & 0 & 0 \end{vmatrix}$$

$$v_{0i} = 4n$$

$$v_{ii} = 8n$$

#### Aim:

make pairwise variances lower than in other designs, whether or not there are period effects.

#### Aim:

make pairwise variances lower than in other designs, whether or not there are period effects.

$$s_{ik} = \begin{cases} \frac{m}{2^k} & \text{if } i = 0\\ \\ \frac{m}{2^{k-i+1}} & \text{if } 0 < i \le k\\ \\ 0 & \text{otherwise.} \end{cases}$$

#### Aim:

make pairwise variances lower than in other designs, whether or not there are period effects.

$$s_{ik} = \begin{cases} \frac{m}{2^k} & \text{if } i = 0 \\ \frac{m}{2^{k-i+1}} & \text{if } 0 < i \le k \end{cases}$$
 Example:  $n = 3, m = 8$ 

$$\frac{m}{2^{k-i+1}} & \text{if } 0 < i \le k \end{cases}$$
 Obose  $\begin{vmatrix} 0 & 1 & 2 & 3 \\ \hline \text{Cohort 1} & 4 & 4 & 0 & 0 \\ \hline \text{Cohort 2} & 2 & 2 & 4 & 0 \\ \hline \text{Cohort 3} & 1 & 1 & 2 & 4 \end{cases}$ 

#### Aim:

make pairwise variances lower than in other designs, whether or not there are period effects.

In Cohort 1:  $\frac{m}{2}$  subjects get dose 1;  $\frac{m}{2}$  subjects get placebo.

In Cohort i:  $\frac{m}{2}$  subjects get dose i; remaining subjects are allocated as in Cohort i-1 with numbers halved  $\mathbb{R}^{2}$ 



			C	oh	ort	1				C	oh	ort	2				C	oh	ort	3		
Dose	0	0	0	0	1	1	1	0	0	1	1	2	2	2	0	1	2	2	3	3	3	3

Dose 0 0 0 0 1 1 1 1 1 0 0 1 1 2 2 2 2 2 0 0 1 2 2 3 3 3 3				C	oh	ort	1				C	oh	ort	2				C	oh	ort	3		
	Dose	1	0	0	0	1	1	1		0	1	1	2	2	2		1	2	2	3	3	3	3

cohort 1 cohort 2 cohort 3 
$$\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$$
 not estimable not estimable estimable

		(	Coh	101	t	1					C	oh	ort	2					C	oh	ort	3		
Dose	0 0	0	0	1		1	1	1	0	0	1	1	2	2	2	2		1	2	2	3	3	3	3
$\overline{Z_3}$																	+	+	+	+	_	_	_	-

cohort 1 cohort 2 cohort 3 
$$\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$$
 not estimable not estimable estimable

 $Z_3$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$ .

			C	oh	ort	1					C	oh	ort	2					C	oh	ort	3		
Dose	0	0	0	0	1	1	1	1	0	0	1	1	2	2	2	2		1	2	2	3	3	3	3
$\overline{Z_3}$																	+	+	+	+	_	_	_	-

cohort 1 cohort 2 cohort 3 
$$\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$$
 not estimable not estimable orthogonally estimable

 $Z_3$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$ .

			C	oh	ort	1					C	oh	ort	2					C	oh	ort	3		
Dose	0	0	0	0	1	1	1	1	0	0	1	1	2	2	2	2	0	1	2	2	3	3	3	3
$Z_3$ $Z_2$									+	+	+	+	_	_	_	_	++	+	+	+	_	_	_	_

 $Z_3$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$ .  $Z_2$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 - 2\tau_2$ .

			C	oh	ort	1					C	oh	ort	2					C	oh	ort	3		
Dose	0	0	0	0	1	1	1	1	0	0	1	1	2	2	2	2	0	1	2	2	3	3	3	3
$Z_3$ $Z_2$									+	+	+	+	_	_	_	_	++	+	+	+	_	_	_	_

 $Z_3$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$ .  $Z_2$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 - 2\tau_2$ .

	Cohort 1	Cohort 2	Cohort 3
Dose	0 0 0 0 1 1 1 1	0 0 1 1 2 2 2 2	0 1 2 2 3 3 3 3
$\overline{Z_3}$			++++
$Z_3$ $Z_2$		++++	++
$Z_1$	++++	++	+ -

 $Z_3$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 + 2\tau_2 - 4\tau_3$ .

 $Z_2$  is the best linear unbiased estimator of  $\tau_0 + \tau_1 - 2\tau_2$ .

 $Z_1$  is the best linear unbiased estimator of  $\tau_0 - \tau_1$ .

# Calculating variances in the halving design (continued)

```
In general, put Z_j = (sum of repsonses on doses 0, ..., j-1 in cohorts j, ..., n) – (sum of responses on dose j)
```

The  $Z_j$  are uncorrelated, with known variances. Linear combinations of them give estimators of all contrasts in the doses.

Hence ...

## Calculating variances in the halving design (continued)

In general, put 
$$Z_j =$$
 (sum of repsonses on doses  $0, ..., j-1$  in cohorts  $j, ..., n$ ) – (sum of responses on dose  $j$ )

The  $Z_j$  are uncorrelated, with known variances. Linear combinations of them give estimators of all contrasts in the doses.

Hence ...

$$v_{ij} = n2^{n} \left( \sum_{t=i}^{j-1} \frac{1}{f(t)} + \frac{4}{f(j)} \right) \quad \text{if } 0 < i < j$$

$$v_{0j} = v_{1j} \quad \text{if } 1 < j$$

$$v_{01} = n2^{n} \frac{4}{f(1)}$$

where

$$f(j) = 2^{n+1} - 2^j.$$



$$n = 3, m = 8$$
  
Numbers of subjects

				J	
				2	
т	Cohort 1	2	6	0	0
1	Cohort 2	2	0	6	0
	Cohort 1 Cohort 2 Cohort 3	2	0	0	6

	n = 3,	m	= 1	8		Scaled variance of	diffei	ences	
	Numbers	of s	sub	jec	ts	r	о со	hort e	ffect
	Dose	0	1	2	3		1	2	3
Т	Cohort 1	2	6	0	0	0	8.0	8.0	8.0
1	Cohort 2	2	0	6	0	1		8.0	8.0
	Cohort 3	2	0	0	6	2			8.0
		'							
	Dose	0	1	2	3		1	2	3
S	Cohort 1	4	4	0	0	$\overline{0}$	8.0	8.0	8.0
S	Cohort 2	4	0	4	0	1		12.0	12.0
	Cohort 3	4	0	0	4	2			12.0
		'					'		
	Dose	0	1	2	3		1	2	3
Н	Cohort 1	4	4	0	0	0	6.9	7.4	9.4
11	Cohort 2	2	2	4	0	1		7.4	9.4
	Cohort 3	1	1	2	4	2			10.0

	n = 3,	m	= :	8		Scaled variance of	diffe	ences	
	Numbers	of s	sub	jec	ts	r	о со	hort e	ffect
	Dose	0	1	2	3		1	2	3
Т	Cohort 1	2	6	0	0	0	8.0	8.0	8.0
1	Cohort 2	2	0	6	0	1		8.0	8.0
	Cohort 3	2	0	0	6	2			8.0
	Dose	0	1	2	3		1	2	3
S	Cohort 1	4	4	0	0	$\overline{0}$	8.0	8.0	8.0
S	Cohort 2	4	0	4	0	1		12.0	12.0
	Cohort 3	4	0	0	4	2			12.0
							'		
	Dose	0	1	2	3		1	2	3
Н	Cohort 1	4	4	0	0	$\overline{0}$	6.9	7.4	9.4
11	Cohort 2	2	2	4	0	1		7.4	9.4
	Cohort 3	1	1	2	4	2			10.0

	n = 3,	m	=	8			Sc	aled v	arianc	e of o	liffer	ences	
	Numbers	of :	sub	jec	ts	fit	ting c	ohort	effect	n	o co	hort e	ffect
	Dose	0	1	2	3		1	2	3		1	2	3
Т	Cohort 1	2	6	0	0	0	16.0	16.0	16.0	0	8.0	8.0	8.0
1	Cohort 2	2	0	6	0	1		32.0	32.0	1		8.0	8.0
	Cohort 3	2	0	0	6	2			32.0	2			8.0
											'		
	Dose	0	1	2	3		1	2	3		1	2	3
S	Cohort 1	4	4	0	0	0	12.0	12.0	12.0	0	8.0	8.0	8.0
3	Cohort 2	4	0	4	0	1		24.0	24.0	1		12.0	12.0
	Cohort 3	4	0	0	4	2			24.0	2			12.0
		'					1				'		
	Dose	0	1	2	3		1	2	3		1	2	3
Н	Cohort 1	4	4	0	0	0	6.9	9.7	15.7	0	6.9	7.4	9.4
п	Cohort 2	2	2	4	0	1		9.7	15.7	1		7.4	9.4
	Cohort 3	1	1	2	4	2			14.0	2			10.0
							'						

	n = 3,	m	=	8			Sc	aled v	arianc	e of o	liffer	ences	
	Numbers	of s	sub	jec	ets	fit	ting c	ohort	effect	n	o co	hort e	ffect
	Dose	0	1	2	3		1	2	3		1	2	3
Т	Cohort 1	2	6	0	0	0	16.0	16.0	16.0	0	8.0	8.0	8.0
1	Cohort 2	2	0	6	0	1		32.0	32.0	1		8.0	8.0
	Cohort 3	2	0	0	6	2			32.0	2			8.0
											'		
	Dose	0	1	2	3		1	2	3		1	2	3
S	Cohort 1	4	4	0	0	0	12.0	12.0	12.0	0	8.0	8.0	8.0
3	Cohort 2	4	0	4	0	1		24.0	24.0	1		12.0	12.0
	Cohort 3	4	0	0	4	2			24.0	2			12.0
		'					1				'		
	Dose	0	1	2	3		1	2	3		1	2	3
Н	Cohort 1	4	4	0	0	0	6.9	9.7	15.7	0	6.9	7.4	9.4
П	Cohort 2	2	2	4	0	1		9.7	15.7	1		7.4	9.4
	Cohort 3	1	1	2	4	2			14.0	2			10.0
							'						

	n = 3,	=	8			Scaled variance of differences								
	Numbers	sub	jec	ts	fit	itting cohort effect				no cohort effect				
T	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	2	6	0	0	0	16.0	16.0	16.0	0	8.0	8.0	8.0	
	Cohort 2	2	0	6	0	1		32.0	32.0	1		8.0	8.0	
	Cohort 3	2	0	0	6	2			32.0	2			8.0	
											'			
S	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	4	4	0	0	0	12.0	12.0	12.0	0	8.0	8.0	8.0	
	Cohort 2	4	0	4	0	1		24.0	24.0	1		12.0	12.0	
	Cohort 3	4	0	0	4	2			24.0	2			12.0	
		'					•				'			
Н	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	4	4	0	0	0	6.9	9.7	15.7	0	6.9	7.4	9.4	
	Cohort 2	2	2	4	0	1		9.7	15.7	1		7.4	9.4	
	Cohort 3	1	1	2	4	2			14.0	2			10.0	

	n = 3,	=	8			Scaled variance of differences								
	Numbers	sub	jec	ts	fit	itting cohort effect				no cohort effect				
T	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	2	6	0	0	0	16.0	16.0	16.0	0	8.0	8.0	8.0	
	Cohort 2	2	0	6	0	1		32.0	32.0	1		8.0	8.0	
	Cohort 3	2	0	0	6	2			32.0	2			8.0	
											'			
S	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	4	4	0	0	0	12.0	12.0	12.0	0	8.0	8.0	8.0	
	Cohort 2	4	0	4	0	1		24.0	24.0	1		12.0	12.0	
	Cohort 3	4	0	0	4	2			24.0	2			12.0	
		'					'				'			
Н	Dose	0	1	2	3		1	2	3		1	2	3	
	Cohort 1	4	4	0	0	0	6.9	9.7	15.7	0	6.9	7.4	9.4	
	Cohort 2	2	2	4	0	1		9.7	15.7	1		7.4	9.4	
	Cohort 3	1	1	2	4	2			14.0	2			10.0	

### How do we calculate variance?

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \mathrm{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \mathrm{information\ matrix} = \mathbf{R} - m^{-1}\Lambda$ 

### How do we calculate variance?

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \mathrm{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \mathrm{information\ matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

### How do we calculate variance?

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \mathrm{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \mathrm{information\ matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

► Find L<sup>−</sup> numerically.

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \operatorname{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \operatorname{information matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

Find L<sup>-</sup> numerically.
 Slightly tricky, because L is not invertible.
 No good for general formulae.

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \operatorname{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \operatorname{information matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

- Find L<sup>-</sup> numerically.
   Slightly tricky, because L is not invertible.
   No good for general formulae.
- Express **L** in spectral form as  $\sum \mu_i \mathbf{P}_i$ ; then  $\mathbf{L}^- = \sum \mu_i^{-1} \mathbf{P}_i$ .

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \operatorname{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \operatorname{information matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

- Find L<sup>-</sup> numerically.
   Slightly tricky, because L is not invertible.
   No good for general formulae.
- Express L in spectral form as  $\sum \mu_i \mathbf{P}_i$ ; then  $\mathbf{L}^- = \sum \mu_i^{-1} \mathbf{P}_i$ . Feasible if the eigenvectors are "nice" and there are few distinct eigenvalues.

$$r_i$$
 = replication of dose  $i = \sum_k s_{ik}$   
 $\lambda_{ij}$  = concurrence of  $i$  and  $j$  in cohorts =  $\sum_k s_{ik} s_{jk}$ 

$$\mathbf{R} = \operatorname{diag}(r_i)$$
  $\Lambda = [\lambda_{ij}]$   $\mathbf{L} = \operatorname{information matrix} = \mathbf{R} - m^{-1}\Lambda$ 

If all responses are uncorrelated with variance  $\sigma^2$  then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \mathbf{L}^{-} \mathbf{x} \sigma^2$ .

- Find L<sup>-</sup> numerically.
   Slightly tricky, because L is not invertible.
   No good for general formulae.
- Express **L** in spectral form as  $\sum \mu_i \mathbf{P}_i$ ; then  $\mathbf{L}^- = \sum \mu_i^{-1} \mathbf{P}_i$ . Feasible if the eigenvectors are "nice" and there are few distinct eigenvalues.
- ► *Ad hoc* methods for special patterns.



### Dose-escalation trials: extended designs

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n+1 cohorts of m subjects each.

#### Dose-escalation trials: extended designs

There are *n* doses, with dose  $1 < \text{dose } 2 < \cdots < \text{dose } n$ .

0 denotes the placebo.

There are n + 1 cohorts of m subjects each.

Cohort 1 subjects may receive only dose 1 or placebo.

In Cohort *i*, for  $2 \le i \le n$ , some subjects receive dose *i*; no subject receives dose *j* if j > i.

In Cohort n + 1, any dose, or placebo, may be used.

### Extended textbook design

Maintain overall equal replication in the final cohort.

$$s_{i,n+1} = \frac{m}{n+1}$$
 for  $i = 0, ..., n$ 

### Extended textbook design

Maintain overall equal replication in the final cohort.

$$s_{i,n+1} = \frac{m}{n+1}$$
 for  $i = 0, ..., n$ 

Example: 
$$n = 4, m = 15$$

Dose	0	1	2	3	4
Cohort 1	3	12	0	0	0
Cohort 2	3	0	12	0	0
Cohort 3	3	0	0	12	0
Cohort 4	3	0	0	0	12
Cohort 5	3	3	3	3	3

### Extended Senn design

In the final cohort, compensate for the previous over-replication of placebo.

$$s_{i,n+1} = \begin{cases} 0 & \text{if } i = 0\\ \frac{m}{n} & \text{otherwise} \end{cases}$$

## Extended Senn design

In the final cohort, compensate for the previous over-replication of placebo.

$$s_{i,n+1} = \begin{cases} 0 & \text{if } i = 0\\ \frac{m}{n} & \text{otherwise} \end{cases}$$

Example: n = 4, m = 16

Dose	0	1	2	3	4
Cohort 1	8	8	0	0	0
Cohort 2	8	0	8	0	0
Cohort 3	8	0	0	8	0
Cohort 4	8	0	0	0	8
Cohort 5	0	4	4	4	4

### Extended halving design

Repeat the final cohort of the standard halving design, to improve comparisons with the highest dose and achieve equal replication overall.

### Extended halving design

Repeat the final cohort of the standard halving design, to improve comparisons with the highest dose and achieve equal replication overall.

Example: 
$$n = 4, m = 16$$

Dose	0	1	2	3	4
Cohort 1	8	8	0	0	0
Cohort 2	4	4	8	0	0
Cohort 3	2	2	4	8	0
Cohort 4	1	1	2	4	8
Cohort 5	1	1	2	4	8

Example: n = 4, m = 16

Example:	n =	4, m	=4
----------	-----	------	----

Dose	0	1	2	3	4
Cohort 1	8	8	0	0	0
Cohort 2	8	0	8	0	0
Cohort 3	8	0	0	8	0
Cohort 4	8	0	0	0	8
Cohort 5	0	4	4	4	4

Dose	0	1	2	3	4
Cohort 1		2	0	0	0
Cohort 2		0	2	0	0
Cohort 3	2	0	0	2	0
Cohort 4	2	0	0	0	2
Cohort 5	0	1	1	1	1

Example: 
$$n = 4, m = 16$$

Dose	0	1	2	3	4
Cohort 1	8	8	0	0	0
Cohort 2	8	0	8	0	0
Cohort 3	8	0	0	8	0
Cohort 4	8	0	0	0	8
Cohort 5	0	4	4	4	4

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

Put  $\mathbf{u} = (4, -1, -1, -1, -1)^{\top} = \text{contrast between dose } 0$  and the rest. Put  $\mathbf{x} = (0, 1, -1, 0, 0)^{\top}$  or any other contrast between non-zero doses.

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

Put  $\mathbf{u} = (4, -1, -1, -1, -1)^{\top} = \text{contrast between dose } 0$  and the rest. Put  $\mathbf{x} = (0, 1, -1, 0, 0)^{\top}$  or any other contrast between non-zero doses. Then  $\mathbf{L}\mathbf{u} = 5\mathbf{u}$  and  $\mathbf{L}\mathbf{x} = 2\mathbf{x}$ ,

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

Put  $\mathbf{u} = (4, -1, -1, -1, -1)^{\top} = \text{contrast between dose } 0$  and the rest. Put  $\mathbf{x} = (0, 1, -1, 0, 0)^{\top}$  or any other contrast between non-zero doses. Then  $\mathbf{L}\mathbf{u} = 5\mathbf{u}$  and  $\mathbf{L}\mathbf{x} = 2\mathbf{x}$ , so

$$\mathbf{L} = 5\mathbf{P}_1 + 2\mathbf{P}_2,$$

where  $\mathbf{P}_1 = \text{ the idempotent for contrast } \mathbf{u}$  and  $\mathbf{P}_2 = \text{ the idempotent for contrasts among the non-zero doses.}$ 

$$4\mathbf{L} = \begin{bmatrix} 16 & -4 & -4 & -4 & -4 \\ -4 & 7 & -1 & -1 & -1 \\ -4 & -1 & 7 & -1 & -1 \\ -4 & -1 & -1 & 7 & -1 \\ -4 & -1 & -1 & -1 & 7 \end{bmatrix}$$

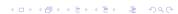
Put  $\mathbf{u} = (4, -1, -1, -1, -1)^{\top} = \text{contrast between dose } 0$  and the rest. Put  $\mathbf{x} = (0, 1, -1, 0, 0)^{\top}$  or any other contrast between non-zero doses. Then  $\mathbf{L}\mathbf{u} = 5\mathbf{u}$  and  $\mathbf{L}\mathbf{x} = 2\mathbf{x}$ , so

$$\mathbf{L} = 5\mathbf{P}_1 + 2\mathbf{P}_2,$$

where  $\mathbf{P}_1=$  the idempotent for contrast  $\mathbf{u}$  and  $\mathbf{P}_2=$  the idempotent for contrasts among the non-zero doses.

$$\mathbf{L}^{-} = \frac{1}{5} \mathbf{P}_{1} + \frac{1}{2} \mathbf{P}_{2},$$

Hence pairwise variances can be calculated.



# Variances in three extended designs

#### Textbook design

#### Senn design

$$v_{0i} = \frac{(n+1)^2(n+2)}{2n+1}$$
  $v_{0j} = \frac{4(n+1)(4+n^2)}{n(4+n)}$   
 $v_{ij} = \frac{2(n+1)^3}{2n+1}$   $v_{ij} = \frac{8n(n+1)}{4+n}$ 

## Variances in three extended designs

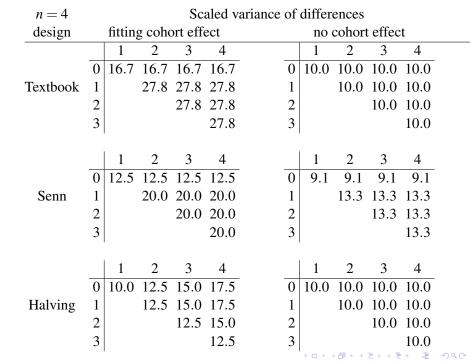
#### Textbook design

#### Senn design

$$v_{0i} = \frac{(n+1)^2(n+2)}{2n+1}$$
  $v_{0j} = \frac{4(n+1)(4+n^2)}{n(4+n)}$   
 $v_{ij} = \frac{2(n+1)^3}{2n+1}$   $v_{ij} = \frac{8n(n+1)}{4+n}$ 

#### Halving design

$$v_{ij} = \frac{(4+j-i)(n+1)}{2}$$
 if  $0 < i < j$   
 $v_{0j} = v_{1j}$  if  $1 < j$   
 $v_{01} = 2(n+1)$ 



n = 4	Scaled variance of differences											
design	fitting cohort effect				ect	av'e		no	cohor	t effec	t	av'e
		1	2	3	4			1	2	3	4	
	0	16.7	16.7	16.7	16.7		0	10.0	10.0	10.0	10.0	
Textbook	1		27.8	27.8	27.8	23.3	1		10.0	10.0	10.0	10.0
	2			27.8	27.8		2			10.0	10.0	
	3				27.8		3				10.0	
		l						ı				
		1	2	3	4			1	2	3	4	
	0	12.5	12.5	12.5	12.5		0	9.1	9.1	9.1	9.1	
Senn	1		20.0	20.0	20.0	17.0	1		13.3	13.3	13.3	11.7
	2			20.0	20.0		2			13.3	13.3	
	3				20.0		3				13.3	
		l						I				
		1	2	3	4			1	2	3	4	
	0	10.0	12.5	15.0	17.5		0	10.0	10.0	10.0	10.0	
Halving	1		12.5	15.0	17.5	14.0	1		10.0	10.0	10.0	10.0
	2			12.5	15.0		2			10.0	10.0	
	3				12.5		3				10.0	
		1						<b>↓</b> □ ▶ ·	( <b>a</b> ) (	≣ > ∢ ≣	→ 重	990

### Average pairwise variance

#### Theorem (Standard)

For a connected design with information matrix  ${\bf L}$ , the average of the pairwise variances is  $2S\sigma^2$ , where S is the average of the reciprocals of the non-zero eigenvalues of  ${\bf L}$ .

#### Random cohort effects

Now assume that the expectation of the response of a subject who gets dose i in cohort k is  $\tau_i$ , and that cohort effects are uncorrelated random variables with common variance  $\sigma_C^2$ .

Put 
$$\mathbf{C}_{\alpha\beta} = \begin{cases} 1 & \text{if subjects } \alpha \text{ and } \beta \text{ are in the same cohort} \\ 0 & \text{otherwise.} \end{cases}$$

Then the variance-covariance matrix of the responses is

$$\sigma^2 \left( \mathbf{I} - \frac{1}{m} \mathbf{C} \right) + \sigma^2 \theta^{-1} \frac{1}{m} \mathbf{C}$$

where 
$$\sigma^2 + m\sigma_C^2 = \theta^{-1}\sigma^2$$
, so  $\theta \in [0, 1]$  with  $\theta = 0$  if cohort effects are fixed  $\theta = 1$  if cohort effects are zero.

If we know the value of  $\theta$  and combine within-cohort and between-cohort information, then the variance of the contrast  $\mathbf{x}^{\top} \boldsymbol{\tau}$  is  $\mathbf{x}^{\top} \left( \mathbf{L} + \theta \widetilde{\mathbf{L}} \right)^{-} \mathbf{x} \sigma^{2}$ , where

$$\mathbf{L} = \operatorname{diag}(r_i) - m^{-1} \Lambda$$
  
$$\widetilde{\mathbf{L}} = m^{-1} \Lambda - \left(\sum r_i\right)^{-1} [r_i r_j].$$

If we know the value of  $\theta$  and combine within-cohort and between-cohort information, then the variance of the contrast  $\mathbf{x}^{\top}\boldsymbol{\tau}$  is  $\mathbf{x}^{\top}\left(\mathbf{L}+\theta\widetilde{\mathbf{L}}\right)^{-}\mathbf{x}\sigma^{2}$ , where

$$\mathbf{L} = \operatorname{diag}(r_i) - m^{-1} \Lambda$$
  

$$\widetilde{\mathbf{L}} = m^{-1} \Lambda - \left(\sum r_i\right)^{-1} [r_i r_j].$$

If L and  $L^-$  have spectral decompositions  $L = \sum \gamma_i P_i$  and  $\widetilde{L} = \sum \delta_i P_i$ 

If we know the value of  $\theta$  and combine within-cohort and between-cohort information, then the variance of the contrast  $\mathbf{x}^{\top}\boldsymbol{\tau}$  is  $\mathbf{x}^{\top}\left(\mathbf{L}+\theta\widetilde{\mathbf{L}}\right)^{-}\mathbf{x}\sigma^{2}$ , where

$$\mathbf{L} = \operatorname{diag}(r_i) - m^{-1} \Lambda$$
  

$$\widetilde{\mathbf{L}} = m^{-1} \Lambda - \left(\sum r_i\right)^{-1} [r_i r_j].$$

If L and  $L^-$  have spectral decompositions  $L = \sum \gamma_i \mathbf{P}_i$  and  $\widetilde{L} = \sum \delta_i \mathbf{P}_i$ 

If we know the value of  $\theta$  and combine within-cohort and between-cohort information, then the variance of the contrast  $\mathbf{x}^{\top}\boldsymbol{\tau}$  is  $\mathbf{x}^{\top}\left(\mathbf{L}+\theta\widetilde{\mathbf{L}}\right)^{-}\mathbf{x}\sigma^{2}$ , where

$$\mathbf{L} = \operatorname{diag}(r_i) - m^{-1} \Lambda$$
  

$$\widetilde{\mathbf{L}} = m^{-1} \Lambda - \left(\sum r_i\right)^{-1} [r_i r_j].$$

If  $\mathbf{L}$  and  $\mathbf{L}^-$  have spectral decompositions  $\mathbf{L} = \sum \gamma_i \mathbf{P}_i$  and  $\widetilde{\mathbf{L}} = \sum \delta_i \mathbf{P}_i$  then the sum of the reciprocals of the eigenvalues of  $\mathbf{L} + \theta \widetilde{\mathbf{L}}$  is

$$\sum \frac{1}{\gamma_i + \theta \delta_i}$$
.

Otherwise, the average variance must be computed numerically for each value of  $\theta$ .



