

A Comparison of Some Methods for Detection of Safety Signals in Randomised Controlled Clinical Trials

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Overview

The main aim of this presentation is to:

- Compare (by way of a simulation) a number of existing approaches for analysing Adverse Events using groupings in clinical trials.
- Discuss the Adverse Event groupings and methods
- Look at a simulation study and the results
- Summary and conclusions

Adverse Events

- Routinely recorded during a trial
- Severity - Common Terminology Criteria for Adverse Events provides a scale from 1–5 (1 = mild,..., 5 = death)
- Time of occurrence and/or duration
- Effect sizes may be small – long follow up / large numbers of patients
- Many different types of Adverse Events – may have multiple hypotheses

Recent Approaches to Analysing Safety Data

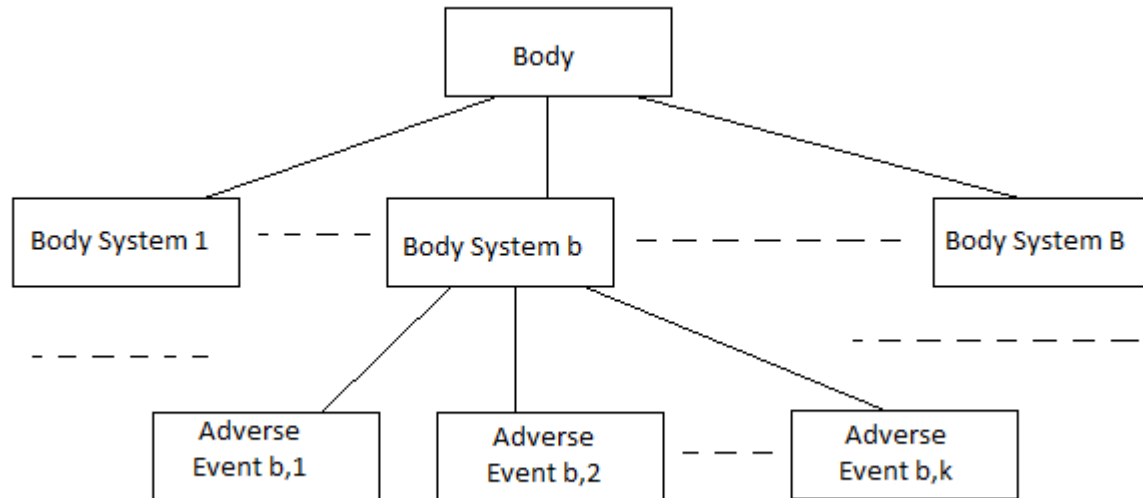
A number of recent approaches to analysing safety data have grouped what they consider to be related adverse events into body-systems or System Organ Classes.

The idea being that if a treatment affects a particular body system then we may expect to see raised adverse event counts for all adverse events in that body-system.

- Berry, Berry - Accounting for Multiplicities in Assessing Drug Safety: A Three-Level Hierarchical Mixture Model (2004)
- Mehrotra, Adewale - Flagging clinical adverse experiences: reducing false discoveries without materially compromising power for detecting true signals (2011).

Body-System Hierarchy

The grouping by Body-System we consider has a natural hierarchical structure, with the body-system being part of an overall body and the adverse events being associated with particular body-systems:



Methods

Method	Description
HIER.BB	Berry and Berry model
HIER.1a	Subset of HIER.BB
BH	Control of the False Discovery Rate by the Benjamini-Hochberg procedure
DFDR	Double False Discovery Rate
NOADJ	Unadjusted testing
BONF	Bonferroni correction
GBH	Group Benjamini-Hochberg
ssBH	Subset Benjamini-Hochberg

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Package: c212 – under development and untested

Berry and Berry Model

B body systems, body-system b containing k_b Adverse Events

N_c – number of patients in the control arm

N_T – number of patients in the treatment arm

X_{bj} – number of adverse events on the control arm

Y_{bj} – number of adverse events on the treatment arm

AE counts: $X_{bj} \sim \text{Bin}(N_c, c_{bj})$, $Y_{bj} \sim \text{Bin}(N_T, t_{bj})$, $1 \leq b \leq B$, $1 \leq j \leq k_b$

Log Odds: $\gamma_{bj} = \text{logit}(c_{bj})$, $\theta_{bj} + \gamma_{bj} = \text{logit}(t_{bj})$

First Level: $\gamma_{bj} \sim N(\mu_{\gamma b}, \sigma_b^2)$, $\theta_{bj} \sim \pi_b I_{[0]} + (1 - \pi_b)N(\mu_{\theta b}, \sigma_b^2)$

θ_{bj} is the log odds-ratio for the occurrence of the AE for treatment compared to control. π_b is the probability that there are no differences in rates in body-system b .

Simulation Study

We used a simulation study to assess how the various methods performed with regard to detecting the raised levels of adverse events between treatment and control.

Model used to generate the simulated trial data:

$$\text{logit}(p_{tbj}) = \mu_{tbj} + U_{tbj}$$

$$X_{bj} \sim \text{Bin}(N_C, p_{1bj})$$

$$Y_{bj} \sim \text{Bin}(N_T, p_{2bj})$$

where μ is a fixed effect and U is a random effect.

Simulation Study

Results from one particular (repeated) simulation:

8 body-systems with between 1 and 11 adverse events in each body-system.
45 adverse events in total.

Trials size:

Trial 1 – 110 patients in each arm

Trial 2 – 450 patients in each arm

Trial 3 – 1100 patients in each arm

For all trials:

The AE rate was raised for body-system 5 for both treatment and control.

The AE rate was raised for body-system 3 for treatment only.

The AE rate was raised for body-system 2 for two out of 4 AEs for treatment only.

Simulation Study

500 simulations in total.

Adverse Event Numbers

In each simulation there are 9 Adverse Events which have underlying rate raised in treatment compared to control.

22500 Adverse Events over the whole simulation.

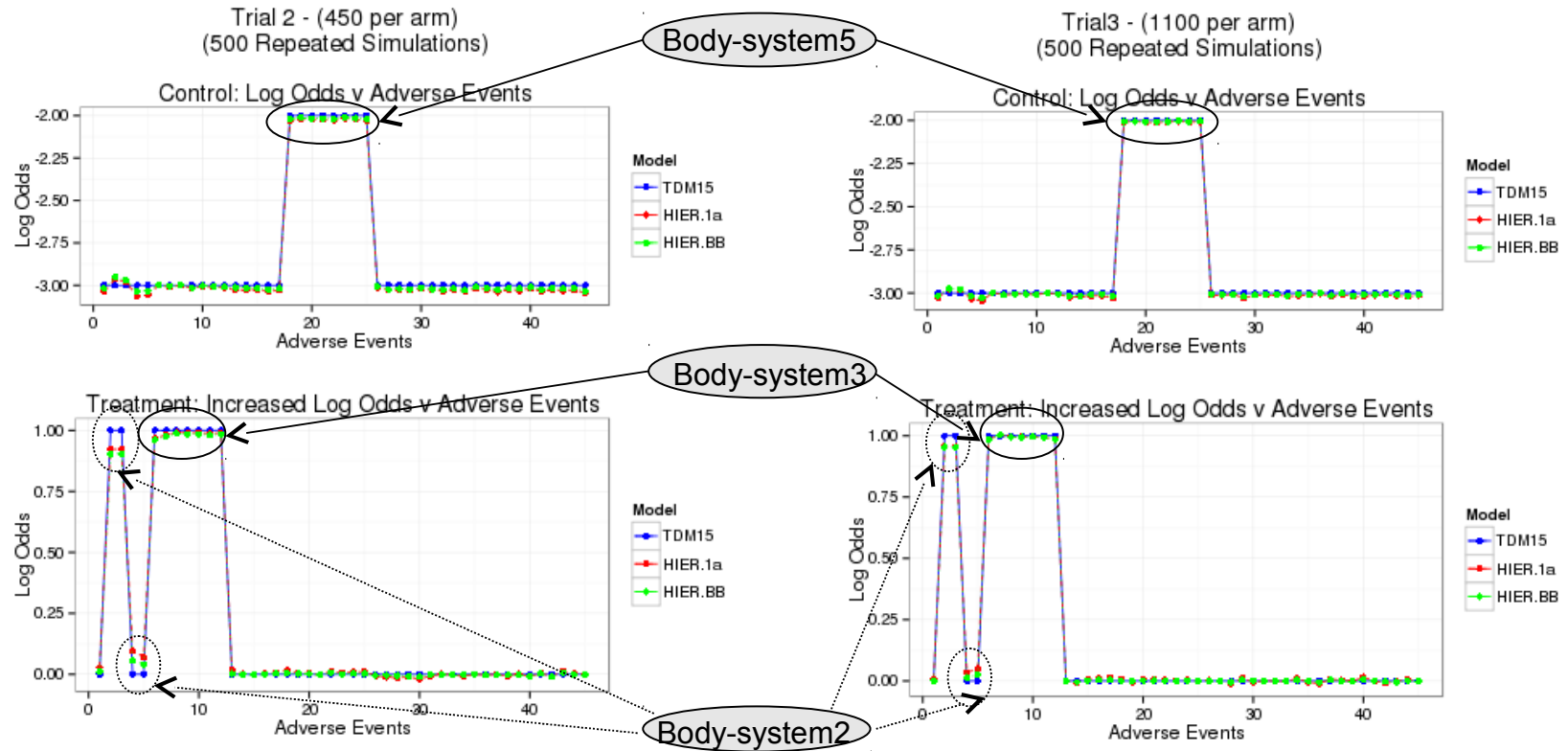
4500 Adverse Events with raised rates over the whole simulation.

Flagging an Adverse Event:

95% posterior probability for Bayesian methods

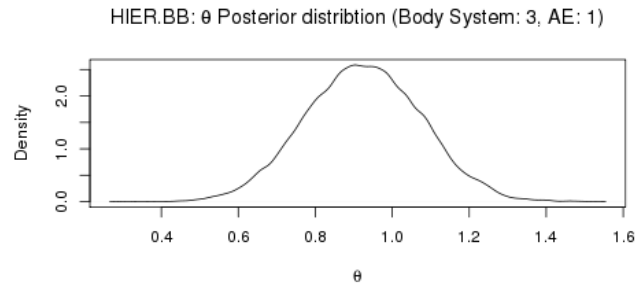
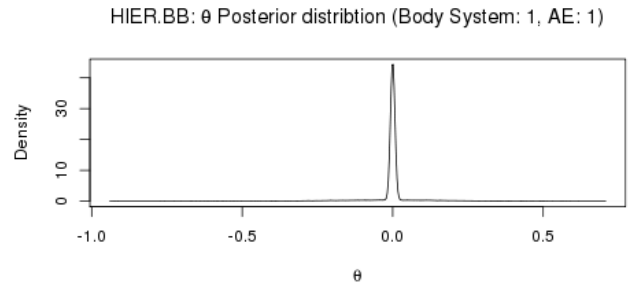
5% significance level for the error controlling methods

Simulation Study

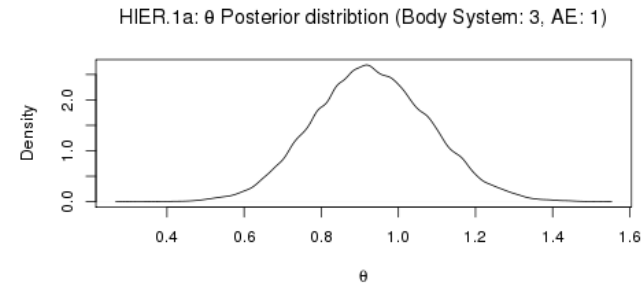
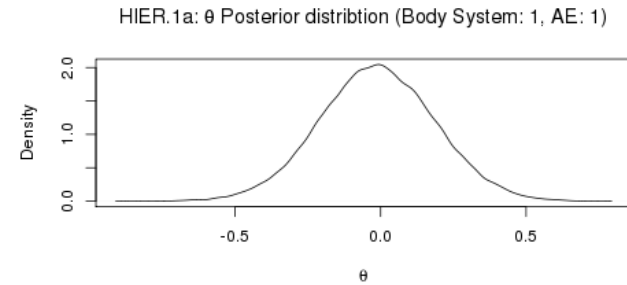


Simulation Study

Berry & Berry Model:
HIER.BB (point mass):



HIER.1a (no point mass):



Simulation Study – Trial 2 (450 per arm)

Method	Correct	Incorrect	Missed
Berry & Berry (HIER.BB)	4303	9	197
Berry & Berry without point mass (HIER.1a)	4492	582	8
Unadjusted Testing (NOADJ)	4374	682	126
Bonferroni Correction (BONF)	3258	12	1242
Double False Discovery Rate (DFDR)	4317	72	183
False Discovery Rate (BH)	4022	114	478
Group Benjamini-Hochberg (GBH)	4441	144	59
Subset Benjamini-Hochberg	3848	14	652

Simulation Study – Trial 3 (1100 per arm)

Method	Correct	Incorrect	Missed
Berry & Berry (HIER.BB)	4498	5	2
Berry & Berry without point mass (HIER.1a)	4500	705	0
Unadjusted Testing (NOADJ)	4500	707	0
Bonferroni Correction (BONF)	4486	10	14
Double False Discovery Rate (DFDR)	4500	67	0
False Discovery Rate (BH)	4499	132	1
Group Benjamini-Hochberg (GBH)	4500	143	0
Subset Benjamini-Hochberg	4498	25	2

Summary

- The simulations have indicated that where there are relationships between the Adverse Events using groupings (body-systems) do appear to make a difference to the results.
- The point mass in the Berry & Berry model (HIER.BB) makes a quantitative difference to the results.
- For the error controlling methods it may be difficult to objectively pick a method of analysing the data before the trial.
- The body-system described in Berry & Berry looks to be a worthwhile structure to consider when modelling data.
- The models and data discussed here do not take into account the severity of events.
- The models and data discussed here do not take into account the timings of events.

References

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