A Group-Sequential Design to Test Efficacy and Inefficiency in Two Subgroups

Darja Tutschkow¹, Astrid Dempfle² & Nina Timmesfeld¹

 ¹Institute of Medical Biometry and Epidemiology, Philipps-University Marburg
² Institute of Medical Informatics and Statistics, Christian Albrechts-University Kiel

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- biomarkers identified as predictive in retrospective or exploratory analyses
- $\bullet\,$ 'issue of multiplicity' $\to\,$ risk of false positive findings
- biomarker-negative subgroup (M^-) not included in later phase III trail \rightarrow no statistically confirmed evidence of inefficiency in M^-

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Example

• trial where the benefit for the M^- was overlooked

- study led by the National Surgical Adjuvant Breast and Bowel Project and the North Central Cancer Treatment Group (Romond et al., 2005)
- effect of Trastuzumab for HER2-positive breast cancer patients
- only HER2-positive patients were included in the trial
- some of initially HER2-positve patients, appeared to be HER2-negative
- subsequently tested HER2-negative patients
- "benefit of adjuvant Trastuzumab may not be limited to patients with HER2 amplification" (Paik, Kim & Wolmark, 2008)

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Group-sequential Design for Both Subgroups

• Assumptions: $X_{Ai}^j \sim \mathcal{N}(\mu_A^j, \sigma^2)$ iid, $X_{Bi}^j \sim \mathcal{N}(\mu_B^j, \sigma^2)$ iid with known σ^2 and $j \in \{+, -\}$

Hypotheses:

$$\begin{split} & H_0^{j,S} : \delta^j \leq 0 \text{ vs. } H_1^{j,S} : \delta^j > 0 \text{ (Superiority)} \\ & H_0^{j,i} : \delta^j \geq \Delta \text{ vs. } H_1^{j,i} : \delta^j < \Delta \text{ (Inefficiency)} \end{split}$$

where $\delta^{j} := \mu_{A} - \mu_{B}$ (difference in treatment effects in subgroup M^{j}) and $\Delta > 0$ (inefficiency margin)

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- predefined number of interim analyses
- equal amount of patients in each subgroup for each interim analysis

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Hierarchical Testing

- first intermin analysis:
 - testing superiority in ${\cal M}^+$ and inefficiency in ${\cal M}^-$
 - if one of either hypotheses is rejected, test it in the other subgroup
- following interim analyses:
 - testing superiority in M^+ and inefficiency in M^- as long as no hypothesis is rejected
 - testing both hypothesis in a subgroup if a hypothesis got rejected in the other subgroup

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Hierarchical Testing - Example



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Simulation

- rejection probabilities for different treatment effects in both subgroups
- expected sample size in both subgroups
- computing the number of required patients N^+ and N^- in M^+ and M^- respectively, such that decision at the last analysis $\rightarrow N^+ = N^- = 2 \cdot 72$
- FWER 5 %
- K=2 analyses
- 10.000 repetitions
- $\Delta = 0.5$
- $\sigma^2 = 1$

Motivation

Results

| δ^+ | δ^{-} | $H_0^{+,S}$ | $H_{0}^{+,I}$ | $H_0^{-,I}$ | $H_0^{-,S}$ | $\mathbb{E}(N^+)$ | $\mathbb{E}(N^{-})$ |
|------------|--------------|-------------|---------------|-------------|-------------|-------------------|---------------------|
| 0 | Δ | 0.025 | 0.024 | 0.025 | 0.024 | 142 | 142 |
| 0 | 2Δ | 0.025 | 0 | 0 | 0.025 | 143 | 143 |
| 0 | 0 | 0.024 | 0.963 | 0.984 | 0 | 100 | 88 |
| Δ | Δ | 0.986 | 0 | 0.025 | 0.962 | 88 | 100 |
| Δ | 0 | 0.979 | 0.021 | 0.980 | 0.020 | 88 | 88 |
| 2Δ | 0 | 1 | 0 | 0.977 | 0.023 | 72 | 87 |
| 2Δ | 2Δ | 1 | 0 | 0 | 1 | 72 | 72 |

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Summary and Outlook

- group-sequantial design to test for superiority and inefficiency for both subgroups for normally distributed data
- next steps:
 - account for different group sizes
 - extension for survival data
- at some point: add more flexibility, e.g.
 - start with the full set, switch to hierarchical procedure and the other way around
 - increase or reduce number of interim analyses
 - change test statistic or outcome measure during the course of the trail, etc. → CRP-method (Müller & Schäfer)

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