



UniversitätsKlinikum Heidelberg

Performance of subgroup selection rules for a targeted therapy in oncology

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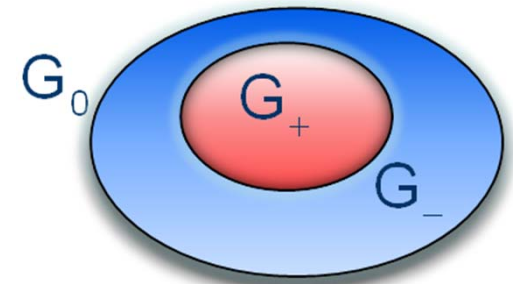
Research developed within the BMBF-funded joint research project **BIMIT**

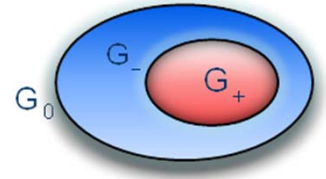




Background

- Total patient population G_0
- Prospectively defined subgroup G_+
 - Potentially increased benefit
 - Identified by biomarker
- Complementary subgroup $G_- := G_0 \setminus G_+$
- Selection of the target population based on data
 - (Pilot / phase II) study A \rightarrow (pivotal / phase III) study B
 - First stage \rightarrow second stage of adaptive enrichment design (e.g. Wang et al. 2007, Jenkins et al. 2011)





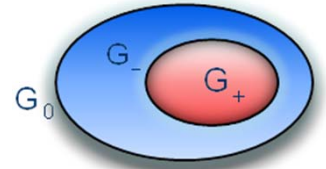
Assumptions and notations

- Assumptions:
 - Binary outcome
 - Two groups: T='treatment', C='control'
- Notations:
 - p_{T+} / p_{T-} : Event probability for patients from subgroup G_+ / G_- assigned to the treatment group
 - p_{C+} / p_{C-} : Event probability for patients from subgroup G_+ / G_- assigned to the control group
 - π : Prevalence of subgroup G_+
 - n : Sample size per group





Basics



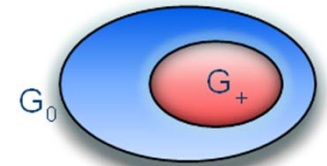
- Effect estimators $\hat{\Delta}_0$ / $\hat{\Delta}_+$ for total population / subgroup:
 - Approximately normally distributed for sufficiently large n
 - $E(\hat{\Delta}_0) = \pi(p_{T+} - p_{C+}) + (1 - \pi)(p_{T-} - p_{C-}) =: \Delta_0$
 - $E(\hat{\Delta}_+) = p_{T+} - p_{C+} =: \Delta_+$
 - $\text{Var}(\hat{\Delta}_0) = [\pi(p_{T+}(1 - p_{T+}) + p_{C+}(1 - p_{C+})) + (1 - \pi)(p_{T-}(1 - p_{T-}) + p_{C-}(1 - p_{C-}))] / n$
 - $\text{Var}(\hat{\Delta}_+) = (p_{T+}(1 - p_{T+}) + p_{C+}(1 - p_{C+})) / (\pi n)$
 - $\text{Cov}(\hat{\Delta}_0, \hat{\Delta}_+) = (p_{T+}(1 - p_{T+}) + p_{C+}(1 - p_{C+})) / n$



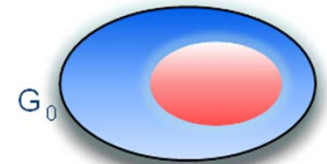
Example for a decision rule

- Rule proposed by Jenkins et al. 2011:

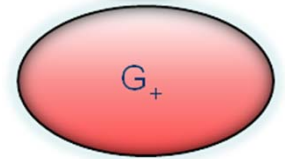
➤ $\hat{\Delta}_0 > c_0, \hat{\Delta}_+ > c_+$: continue with G_0 and G_+



➤ $\hat{\Delta}_0 > c_0, \hat{\Delta}_+ \leq c_+$: continue with G_0 only



➤ $\hat{\Delta}_0 \leq c_0, \hat{\Delta}_+ > c_+$: continue with G_+ only



➤ $\hat{\Delta}_0 \leq c_0, \hat{\Delta}_+ \leq c_+$: stop for futility



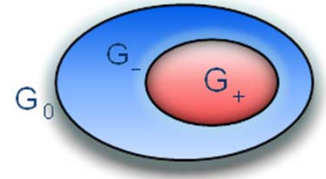
How to choose (c_0, c_+) in case of uncertain $p_{T+} / p_{C+} / p_{T-} / p_{C-}$?

Jenkins M, Stone A, Jennison C (2011). An adaptive seamless phase II/III design for oncology trials with subpopulation selection using correlated survival endpoints. *Pharmaceutical Statistics* 10: 347-356.





Optimal decision rules

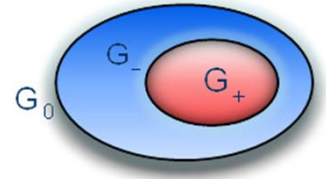


- Assumptions:
 - Selection of G_0 desired if $\Delta_0 > \tau_0$
 - Selection of G_+ desired if $\Delta_+ > \tau_+$
 - *A priori* distributions for $p_{T+} / p_{C+} / p_{T-} / p_{C-}$
 - Quadratic loss function for wrong decision
 - Optimal decision rule minimizes expected loss (“risk”)
- Optimal decision thresholds (c_0^* , c_+^*) can be computed using numerical integration assuming, e.g., continuous uniform priors.





Optimal decision rules (2)



- In the following, we will consider
 - Relevance thresholds $\tau_0 = 0.05$, $\tau_+ = 0.1$
 - Prevalences $\pi = 0.1, 0.25, 0.5$
 - Biomarker assumed to be predictive, i.e.

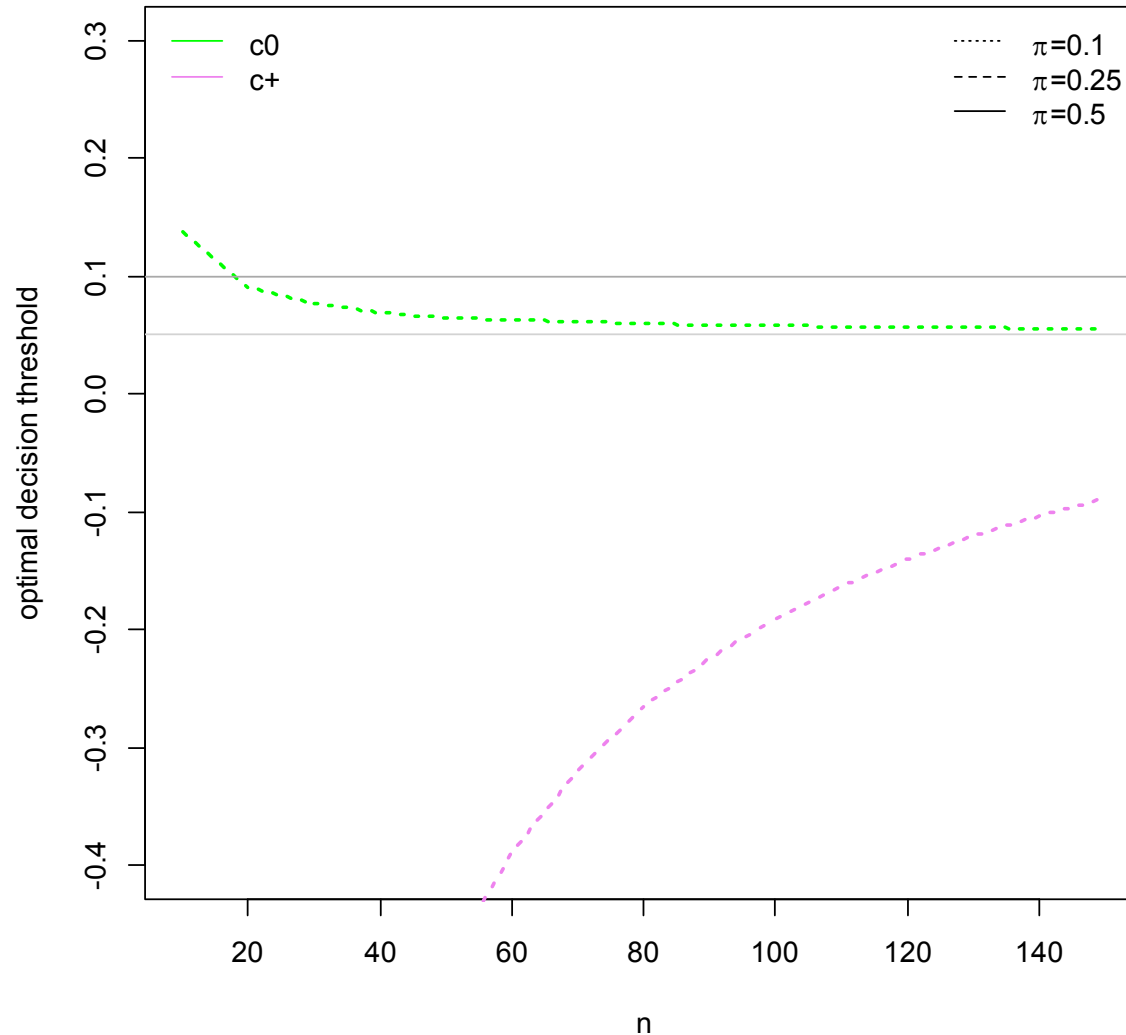
$$p_{T+} \sim U_{[0.3, 0.6]}, \quad p_{T-}, p_{C+}, p_{C-} \sim U_{[0.1, 0.4]}$$





Optimal decision rules (3)

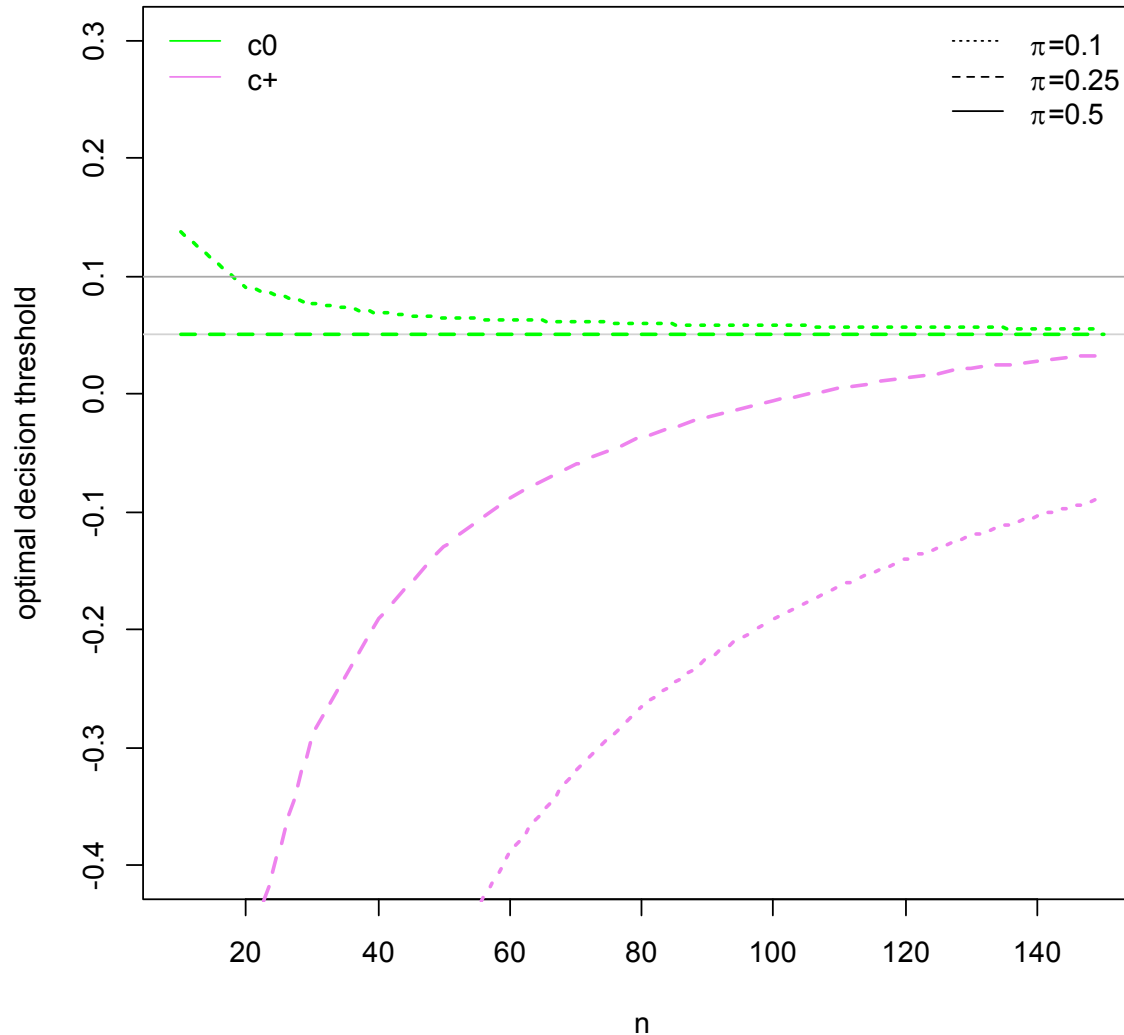
- Example: Optimal decision thresholds for „predictive“ prior





Optimal decision rules (3)

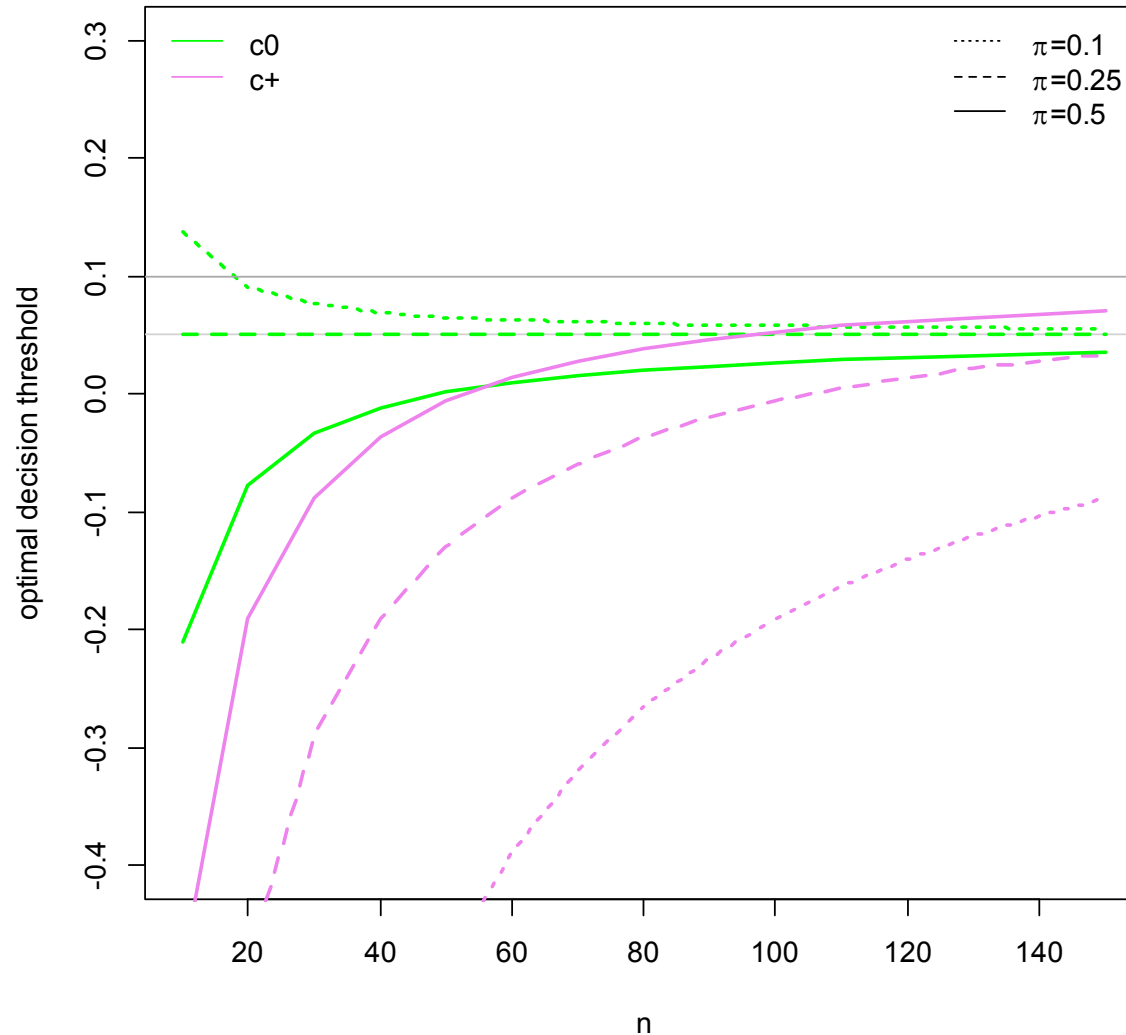
- Example: Optimal decision thresholds for „predictive“ prior





Optimal decision rules (3)

- Example: Optimal decision thresholds for „predictive“ prior





Performance of optimal decision rules – clinical trial example

- Can the incorporation of an optimal decision rule lead to an increase in the probability of the selection of the correct target population?
- Demonstration by clinical trial example:
 - The monoclonal antibody trastuzumab has been shown to be effective as an add-on-therapy exclusively for breast cancer patients which overexpress HER2, which is the case for ~ 20-30% of all patients.
 - However, there is some evidence that trastuzumab might also be effective for HER2-negative patients, which is currently investigated within a large RCT (ClinicalTrials.gov Identifier: NCT01548677).





Assumptions for clinical trial example

- Assume that it is planned to investigate the efficacy of trastuzumab within an enrichment design with subgroup selection at interim with
 - Primary endpoint “event-free survival at five years” (an event is either disease recurrence, progression, or death)
 - $n = 400$, $\pi = 0.2$, $\tau_0 = 0.08$, $\tau_+ = 0.1$
 - Actual event rates
 - $p_{T-} = 0.65$ (Scenario A), $p_{T-} = 0.7$ (Scenario B)
 - $p_{T+} = 0.6$
 - $p_{C-} = 0.6$
 - $p_{C+} = 0.45$





Clinical trial example - *ad hoc* decision rule (a)

- Relevant treatment effect in the subgroup ($\Delta_+ = 0.15 > 0.1 = \tau_+$)
- For Scenario A, i.e. $p_{T-} = 0.65$,
 - $\Delta_0 = 0.07 \leq 0.08 = \tau_0$
 - Selection of G_+ only is desirable
- For Scenario B, i.e. $p_{T-} = 0.7$,
 - $\Delta_0 = 0.11 > 0.08 = \tau_0$
 - Selection of both G_0 and G_+ is desirable
- One could, e.g., simply choose an *ad hoc* decision rule

$(c_0, c_+) = (\tau_0, \tau_+) = (0.08, 0.1) \rightarrow$ **Decision rule (a)**





Clinical trial example – optimal decision rule based on a “predictive” prior (b)

- However, consider the situation of prior knowledge about event rates from a previously conducted trial.
 - Reported 95%-confidence intervals can be used to derive an optimal decision threshold by modelling uniform prior distributions from them.
 - Example (based on results from Gianni et al., 2014):
 - $p_{T+} \sim U_{[0.48, 0.66]}$, $p_{C+} \sim U_{[0.34, 0.52]}$
 - $p_{T-} \sim U_{[0.5, 0.7]}$, $p_{C-} \sim U_{[0.5, 0.7]}$
- Incorporating this prior knowledge yields optimal decision thresholds $(c_0^*, c_+^*) = (0.0822, 0.0601)$ → **Decision rule (b)**



Gianni L et al. (2014). Neoadjuvant and adjuvant trastuzumab in patients with HER2-positive locally advanced breast cancer (NOAH): follow-up of a randomised controlled superiority trial with a parallel HER2-negative cohort. *The Lancet Oncology* 15: 640-647.





Clinical trial example – optimal decision rule based on a non-informative prior (c)

- Alternatively, one could consider non-informative priors to derive optimal decision thresholds, i.e. $p_{T+}, p_{T-}, p_{C+}, p_{C-} \sim U_{[0, 1]}$.
- In this case, the optimal decision rule is

$$(c_0^*, c_+^*) = (0.0807, 0.1029) \rightarrow \text{Decision rule (c)}$$

- We evaluated the performance of decision rules (a), (b), (c) in terms of selection probabilities by Monte-Carlo simulation (1,000,000 simulated studies per scenario)





Selection probabilities for scenario A ($p_{T-} = 0.65$)

decision rule	(a) $(c_0, c_+) =$ $(0.08, 0.1)$	(b) $(c_0^*, c_+^*) =$ $(0.0822, 0.0601)$	(c) $(c_0^*, c_+^*) =$ $(0.0807, 0.1029)$
probability			
select G_0 and G_+	0.3226		
select G_0 only	0.0493		
select G_+ only	0.3919		
stop for futility	0.2361		





Selection probabilities for scenario A ($p_{T-} = 0.65$)

decision rule	(a) $(c_0, c_+) =$ $(0.08, 0.1)$	(b) $(c_0^*, c_+^*) =$ $(0.0822, 0.0601)$	(c) $(c_0^*, c_+^*) =$ $(0.0807, 0.1029)$
probability			
select G_0 and G_+	0.3226	0.3587	
select G_0 only	0.0493	0.0132	
select G_+ only	0.3919	0.5262	
stop for futility	0.2361	0.1077	





Selection probabilities for scenario A ($p_{T-} = 0.65$)

decision rule	(a)	(b)	(c)
probability	$(c_0, c_+) =$ (0.08, 0.1)	$(c_0^*, c_+^*) =$ (0.0822, 0.0601)	$(c_0^*, c_+^*) =$ (0.0807, 0.1029)
select G_0 and G_+	0.3226	0.3587	0.3226
select G_0 only	0.0493	0.0132	0.0493
select G_+ only	0.3919	0.5262	0.3919
stop for futility	0.2361	0.1077	0.2361

- Decision rules (a) and (c) perform identically.
- Relatively high probability for a futility stop for rules (a) and (c).
- Decision rule (b) shows the highest probability for a correct selection.



Selection probabilities for scenario B ($p_{T-} = 0.7$)

decision rule	(a)	(b)	(c)
probability	$(c_0, c_+) =$ (0.08, 0.1)	$(c_0^*, c_+^*) =$ (0.0822, 0.0601)	$(c_0^*, c_+^*) =$ (0.0807, 0.1029)
select G_0 and G_+	0.6232		
select G_0 only	0.1796		
select G_+ only	0.0914		
stop for futility	0.1059		





Selection probabilities for scenario B ($p_{T-} = 0.7$)

decision rule	(a) $(c_0, c_+) =$ $(0.08, 0.1)$	(b) $(c_0^*, c_+^*) =$ $(0.0822, 0.0601)$	(c) $(c_0^*, c_+^*) =$ $(0.0807, 0.1029)$
probability			
select G_0 and G_+	0.6232	0.7419	
select G_0 only	0.1796	0.0609	
select G_+ only	0.0914	0.1431	
stop for futility	0.1059	0.0542	





Selection probabilities for scenario B ($p_{T-} = 0.7$)

decision rule	(a)	(b)	(c)
probability	$(c_0, c_+) =$ (0.08, 0.1)	$(c_0^*, c_+^*) =$ (0.0822, 0.0601)	$(c_0^*, c_+^*) =$ (0.0807, 0.1029)
select G_0 and G_+	0.6232	0.7419	0.6232
select G_0 only	0.1796	0.0609	0.1796
select G_+ only	0.0914	0.1431	0.0914
stop for futility	0.1059	0.0542	0.1059

- Again, decision rules (a) and (c) perform identically.
- As before, decision rule (b) shows the highest probability for a correct selection.



Conclusions

- Optimal decision rules incorporate various aspects of the design of a clinical trial with subgroup selection.
- Under the situation of no prior knowledge on treatment effects, optimal decision rules can also be derived and show a performance comparable to *ad hoc* rules in terms of correct selection probability.
- If there is prior knowledge on treatment effects, optimal decision rules may lead to an increased probability for a correct selection of the target population, which also yields an increased statistical power (Krisam & Kieser, 2015b).





References

- Gianni L et al. (2014).** Neoadjuvant and adjuvant trastuzumab in patients with HER2-positive locally advanced breast cancer (NOAH): follow-up of a randomised controlled superiority trial with a parallel HER2-negative cohort. *The Lancet Oncology* **15**: 640-647.
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- Wang SJ, O'Neill RT, Hung HMJ (2007).** Approaches to evaluation of treatment effect in randomized clinical trials with genomic subset. *Pharmaceutical Statistics* **6**: 227-244.

