



## Blinded Sample Size Re-estimation for Adaptive Enrichment Designs with Longitudinal Data

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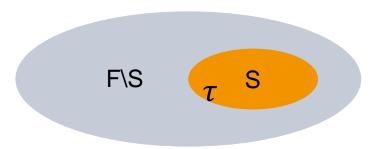
#### **Outline**



- Motivation
- ➤ Adaptive Enrichment Design
- Statistical Model
- ➤ Blinded Sample Size Re-estimation
- > Simulations
- > Discussion
- > References

#### **Motivation**



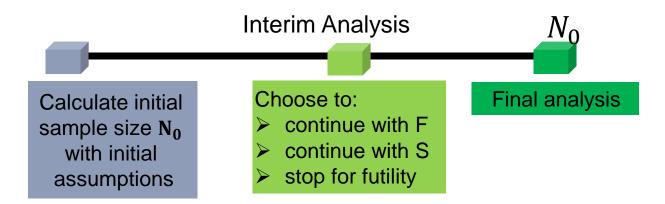


- One major goal in personalized/stratified medicine is the identification of subgroups (S)
- ➤ These subgroups might yield higher efficacy or provide a better safety profile
- ➤ A requirement for adaptive enrichment designs are identifiable subgroups within the population of interest (F) by e.g. genetic markers



### Adaptive Enrichment Design (AED)

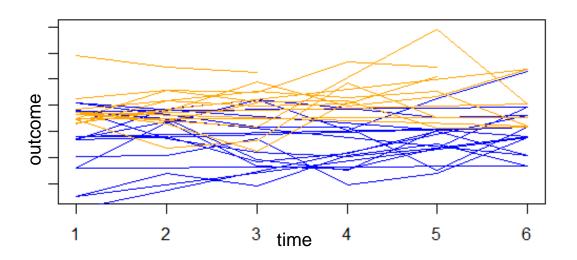
- > Step 1: calculate sample size based on initial assumptions about nuisance parameters and recruit subjects from full-population
- Interim Analysis: based on a decision rule choose to...
  - ...continue study with...
    - ... full-population
    - ... sub-population (enrichment)
  - ...stop study for futility
  - ...based on collected data and a pre-defined selection rule
- Final analysis: test for efficacy using combination tests





### **Datatype**

> Repeated measurements



> In this setting: missingness based on MCAR dropouts

Pat ID	Time					
	t <sub>0</sub>	$t_1$	$t_2$	$t_3$	t <sub>4</sub>	$t_5$
1	2.4	2.8	-	-	-	-
2	2.6	2.8	3.0	3.3	3.5	3.6
3	2.8	3.1	3.1	3.2	-	-
4	2.5	2.8	2.9	3.1	3.3	3.4

#### Statistical Model



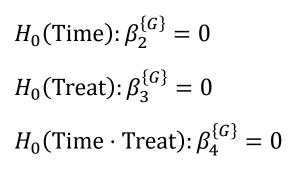
Estimation of linear trends for repeated measures

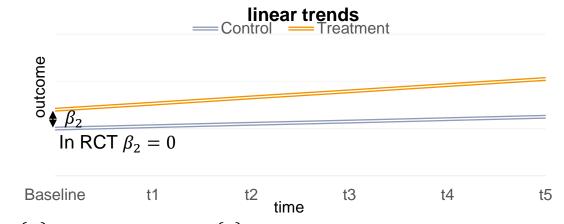
$$y_{ijk}^{\{G\}} = \beta_1^{\{G\}} + \beta_2^{\{G\}} \cdot j + \beta_3^{\{G\}} \cdot 1_{\{k = \text{Treat}\}} + \beta_4^{\{G\}} \cdot j \cdot 1_{\{k = \text{Treat}\}} + \epsilon_{ijk}^{\{G\}}$$

$$\operatorname{cor}(y_{ijk}, y_{ihk}) = \rho^{|j-h|}; \operatorname{cor}(y_{ijk}, y_{oj'k'}) = 0; \ \epsilon_{ijk} \sim N(0, \sigma^2)$$

$$i = 1, \dots, n; j = 0, \dots, t; \ k \in \{\text{Treat}; \text{Contr}\}; G \in \{F; S\}; N = 2 \cdot n$$

Possible hypotheses





 $\succ$  test null hypotheses  $H_0: \beta_4^{\{F\}} \le 0$  and  $H_0: \beta_4^{\{S\}} \le 0$  in a co-primary analysis, controlling the FWER using closed testing procedure [1]



#### **Construction of Test Statistics**

- Let **Σ** denote the covariance matrix of  $\sqrt{n}(\hat{\beta} \beta)$  with  $\Sigma(4,4) = Var(\hat{\beta}_4)^{[2,3,4]}$
- $\Sigma$  depends on model specific parameters:  $\sigma^{\{G\}^2}$ ,  $\rho$ , overall dropout, t
- $\triangleright$  Consider the normalized test statistics of  $\hat{\beta}_4$  for F and S

$$Z^{\{F\}} = \frac{\sqrt{N} \cdot \hat{\beta}_4^{\{F\}}}{\sqrt{\Sigma^{\{F\}}(4,4)}}, \qquad Z^{\{S\}} = \frac{\sqrt{N\tau} \cdot \hat{\beta}_4^{\{S\}}}{\sqrt{\Sigma^{\{S\}}(4,4)}}$$

Test intersection hypothesis assuming a bivariate normal distribution of  $(Z^{\{F\}}, Z^{\{S\}})^{'}$  [5]

$$F_{H_0} = \begin{pmatrix} Z^{\{F\}} \\ Z^{\{S\}} \end{pmatrix}^{H_0: \beta_4^{\{F \cap S\}}} \begin{pmatrix} \begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} 1 & \sqrt{\tau} \\ \sqrt{\tau} & 1 \end{pmatrix} \end{pmatrix}$$

<sup>[3]</sup> Jung and Ahn (2003)

<sup>[4]</sup> Wachtlin and Kieser (2013)



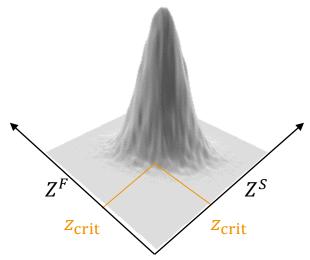
### Sample Size Estimation

ightharpoonup Distribution under  $H_1: \beta_4^{\{G\}} \ge 0$ 

$$F_{H_1} = \begin{pmatrix} Z^{\{F\}} \\ Z^{\{S\}} \end{pmatrix} \approx N \left( \begin{pmatrix} Z^F \\ Z^S \end{pmatrix} \middle| \begin{pmatrix} 1 & \sqrt{\tau} \\ \sqrt{\tau} & 1 \end{pmatrix} \right)$$

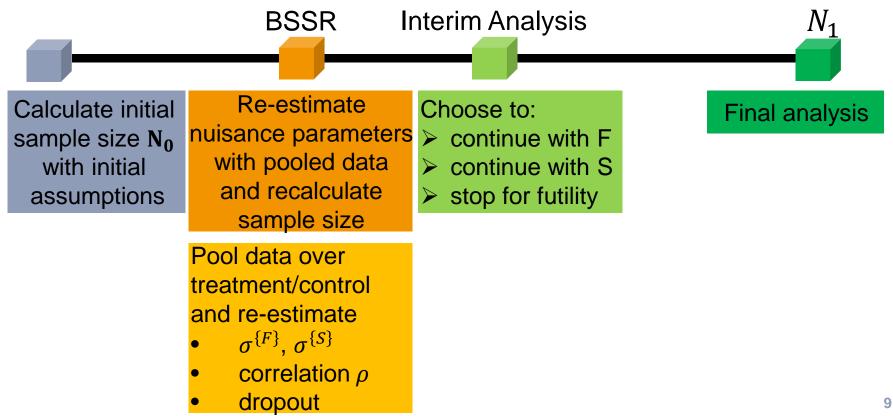
- ightharpoonup Let  $z_{\mathrm{crit}}$  denote the equicoordinated  $\left(1-\frac{\alpha}{2}\right)$  -quantile under  $H_0$
- $\triangleright$  Calculate required sample size iteratively based on  $F_{H_1}$  given  $z_{\text{crit}}$

$$N = \min\{n \mid 1 - F_{H_1}(z_{\text{crit}}) \ge Power\}$$





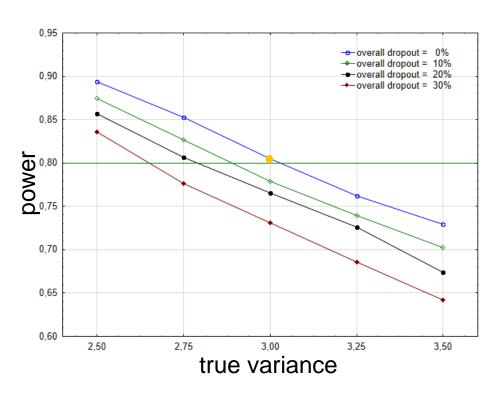
### AED combined with Blinded Sample Size Re-estimation





### Fixed Sample Size Design

Simulations	10,000
$\alpha$	0.025
Power	8.0
$eta_4^{\{F\}} = eta_4^{\{S\}}$	0.1
$\sigma_{F \setminus S}^{2^{ ext{init}}} = \sigma_{S}^{2^{ ext{init}}}$	3
au	0.5
Overall dropout <sub>init</sub>	0%
$N_0$	744



> "The spirit behind **internal pilots** is simple: one uses patients in the pilot to alter the main study, but one does not discard those data from those patients. " <sup>[6]</sup>

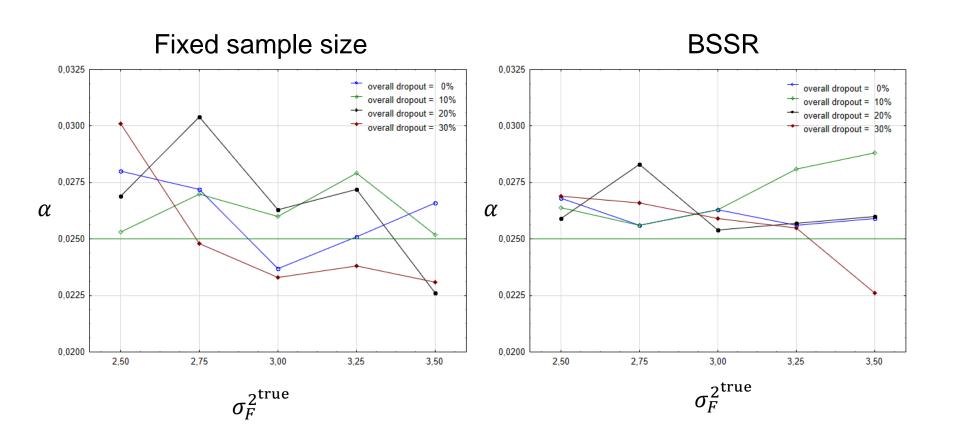


#### Standard values for simulation studies

Simulations	10,000		
$\alpha$	0.025		
Power	0.8		
$ \beta_4^{\{F\}} = \beta_4^{\{S\}} $	0.1		
$\sigma_{F\backslash S}^{2^{\mathrm{init}}} = \sigma_{S}^{2^{\mathrm{init}}}$	3		
$\sigma_{E\backslash S}^{ ext{2true}}$	3		
$\sigma_{ m S}^{ m 2true}$	(2.0, 2.5, 3.0, 3.5, 4.0)		
τ	0.5		
Overall dropout <sub>init</sub>	0%		
Allocation ratio	1:1		
BSSR at	$0.4 \cdot N_0$		
Interim at	$\max(0.4 \cdot N_0, 0.5 \cdot N_1)$		
$N_0$	744		

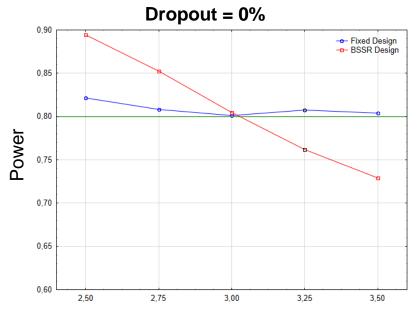


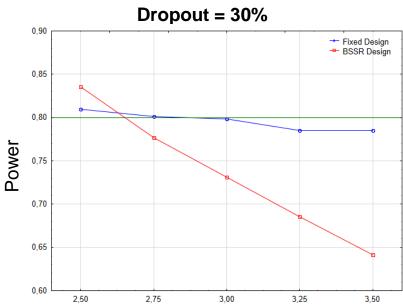
### Type-I-error rate

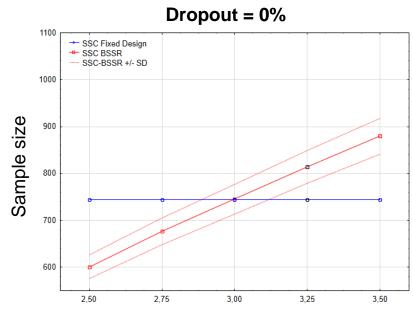


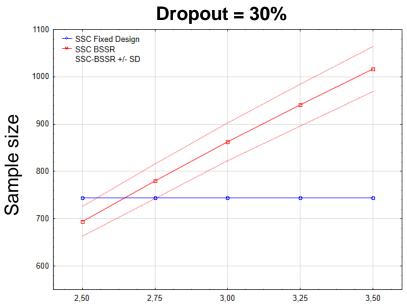


#### Combination of BSSR and AED











#### Discussion & Outlook

- Combination of adaptive enrichment designs and blinded samples size re-estimation provides flexible and robust designs
- Adaptive enrichment design controls type-I-error rate
- BSSR compensates for initially miss-specified nuisance parameters and dropouts in terms of power
- Further investigate and compare weighted-GEE and MI methods in MAR situations
- Extend sample size estimation for cases where sub-population was selected in interim



### For further reading

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- Stallard N. et al. (2013) Adaptive Designs for Confirmatory Clinical Trials with Subgroup Selection. Journal of Biopharmaceutical Statistics



### For further reading

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- Marcus R, Peritz E, Gabriel K. R. (1976) On closed testing procedures with special references to order analysis of variance. Biometrika 63(3):655-660



# Thank you for your attention!