



Methodological Challenges in PrecISE

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- PrecISE (**P**recision **I**nterventions for **S**Evere asthma)
- Collaborative Studies Coordinating Center (CSCC) at the Department of Biostatistics UNC at Chapel Hill was awarded \$61 million by the National Heart, Lung and Blood Institute in September 2017



NHLBI stated the following objectives for PrecISE

1. Run a controlled clinical trial **to evaluate efficacy** of several novel interventions in severe asthma patients
2. Use **precision medicine** approaches
3. Trial design needs to be **adaptive**

Precision medicine approaches in severe asthma

Several asthma treatment recently approved by the FDA:

Treatment	FDA approval	Approved for
FASENRA (benralizumab)	2017	Eosinophilic phenotype, blood eosinophils ≥ 300 cells/ μ l
DUPIXENT (dupilumab)	2018	Eosinophilic phenotype, blood eosinophils ≥ 300 cells/ μ l
Tezspire (tezepelumab)	2021	Unselected population, but better treatment effect was seen in patients with blood eosinophils ≥ 300 cells/ μ l



Interventions that are being evaluated in PrecISE in patients with severe asthma

- **Imatinib** (brand name Gleevec) is an oral chemotherapy to treat patients with cancer
- **Clazakizumab**, a novel therapy currently being investigated in psoriatic arthritis. Not yet approved for any indication
- **Cavosonstat**, a novel treatment, was investigated in cystic fibrosis but did not demonstrate benefit
- **Broncho-Vaxom** is known to support respiratory tract resistance to bacterial infections, has been used in Europe for the last two decades
- **Medium chain triglycerides** (MCT), a food supplement

Biomarker positive subgroups

Intervention	Subgroup	Prevalence
Imatinib	Eos < 300	62%
Clazakizumab	IL-6 > 3.1	33%
Cavosonstat	Genotypes	64%
Broncho-Vaxom	Eos ≥ 300	38%
MCT	FeNO ≥ 15 ppb	64%

Eos = blood eosinophils count

IL-6 = interleukin 6

FeNO = fractional exhaled nitric oxide

PrecISE is a multi-period crossover trial



Option 1: N of 1



**Option 2: a sequence of
2-period crossovers**



Option 3: one random placebo

16-week long periods with 4-week washouts

2 - 6 treatment periods for each participant depending on the time of entry to the study₆

PrecISE is a multi-period crossover trial



Option 1: N of 1



Option 2: a sequence of 2-period crossovers



Option 3: one random placebo



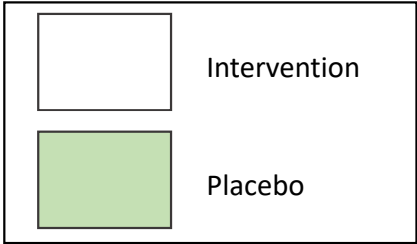
PrecISE: 2-period crossover followed by a sequence of active treatments with possibly 1 more placebo



	Period 1	Period 2	Period 3	Period 4	Period 5
Patient 1		<i>placebo</i>			
Patient 2	<i>placebo</i>		<i>placebo</i>		
Patient 3	<i>placebo</i>				
Patient 4	<i>placebo</i>			<i>placebo</i>	
Patient 5	<i>placebo</i>				
	Period 1	Period 2	Period 3	Period 4	Period 5
	In the first two periods, patients receive the same intervention in active and placebo forms (random sequence).		In subsequent treatment periods, patients will be randomized to different interventions. Participants may also be randomized to receive placebo in periods 3-6, such that on average, 20% of participants will receive a 2nd placebo.		

Other design decisions

- An inclusion of active control
 - Decided not to include
- Number of primary endpoints
 - 3
- Uncertainty about the value of within subject correlation
 - Interim analysis to estimate correlation



Study design in each arm

- Proportion of biomarker positive (A+) and negative participants (A-)
- Test in A+ only? in A-? in unselected?

Recent clinical trials in severe asthma

Treatment	Clinical Trials	Trial Design	Primary and Secondary Analysis
FASENRA (benralizumab)	CALIMA, SIROCCO	66% with blood eos ≥ 300 *	Primary analysis in blood eos ≥ 300
DUPIXENT (dupilumab)	LIBERTY ASTHMA QUEST	44% with blood eos ≥ 300 *	Primary analysis in unselected Prespecified subgroup analyses in blood eos ≥ 300
Tezspire (tezepelumab)	NAVIGATOR	50% with blood eos ≥ 300 *	Primary analysis in unselected Prespecified subgroup analyses in blood eos ≥ 300

*Population prevalence of eosinophilic phenotype (blood eos ≥ 300) in severe asthma is 38%

How to design a trial with a biomarker defined subgroup?

- Phase 2 approach to designing a trial with a subgroup:
 - The goal is to show that the treatment effect is significantly different from 0 in a biomarker negative subgroup (**A-**) and/or a biomarker positive (**A+**)
 - The easiest is to run 2 parallel trials: in A- and A+
 - More efficient options are available (Freidlin et al., 2013; Parashar et al., 2016)
- Phase 3 approach to designing a trial with a subgroup:
 - The goal is to show that the treatment effect is significantly different from 0 in **unselected population** (A- and A+ combined) and/or in **A+**
 - The most efficient way is run a trial in A+ only
 - However, it is often desirable to enroll participants **according to population prevalence** (Rosenblum and Qian, 2016; Rosenblum et al., 2016, Dmitrienko et al., 2017)

Phase 2 versus phase 3 approach

- Phase 2 enrolls to A- and A+ according to the required allocation proportion
- Phase 3 enrolls according to the population prevalence
- **Phase 3 approach requires 20-50% less participants**
 - even more participants need to be screened to find the required number of A- and A+ participants
- When A+ prevalence is high, Phase 3 approach is likely to **conclude that the treatment is effective in unselected population** when only A+ shows activity (Rothmann et al., 2012)

Approaches we considered to design PrecISE

- Enroll A+ only
 - Advantage: will be able to utilize study resources in the most efficient way
 - Disadvantage: This design is **not responsive** to the Request for Application (RFA) since there is no precision medicine component
- Enroll A- and A+ according to the population prevalence and test for treatment effect in unselected and in A+ (Phase 3 approach)
 - Advantage: responsive to the RFA since we can update the biomarker cut-off during the trial (if we can halt enrollment to an intervention) or in a post-hoc analysis
 - Disadvantage: **Not enough power** for interventions with small subgroup

PrecISE Study Design

- Test for efficacy in A+ only
 - No testing for treatment effect in A- or in unselected population
- Enroll more participants from A+ than A-, **2:1 ratio A+/A-**, this is to update the biomarker cut-off (precision medicine component)
 - If the biomarker cut-off is re-estimated during the trial, test for treatment effect in the combined sample of old A+ (before cut-off re-estimation) and a new A+ (after cut-off re-estimation)
 - Imatinib, cavosonstat, and MCT have subgroup prevalence of 64%, enrolling according to population prevalence (as in LIBERTY ASTHMA QUEST trial)
 - Clazakizumab and Broncho-Vaxom have subgroup prevalence of 38% and 33%, need to oversample A+ (as in CALIMA and SIROCCO trials)

How to define the best subgroup?

- Definition 1. The best subgroup is defined as the largest subgroup with a **treatment effect of at least Δ**
- Definition 2. The best subgroup is defined as the subgroup **maximizing**
 $U = \text{Treatment effect} \times \text{Prevalence}^\gamma$
 - When $\gamma = 0$, the treatment effect is maximized
 - When $\gamma = 0.5$, the power is maximized
- In PrecISE, we use **$\gamma = 0.5$** due to the new cut-off being applied prospectively to baseline data of participants already on the treatment

How to update the cut-off at the interim and final analysis?

When updating a cut-off of a single biomarker, a non-parametric approach performed the best (Joshi et al., 2019)

Non-parametric approach : select the subgroup that maximizes

$$U = \text{Treatment effect} \times \text{Prevalence}^\gamma$$

When $\gamma = 0.5$, select the subgroup with **the largest test statistic**

How to adjust for multiplicity in post-hoc subgroup analysis?

- Cross-validation and bootstrap (Simon, 2008; Zhang, et al., 2017)
- Bootstrap (Guo et al., 2020)

PrecISE

- First participant screened Dec 2019
- First participants randomized Aug 2020
- 136 participants have been randomized as of April 1, 2022
- Target sample size 500

References

1. Dmitrienko, A, Millen, B, Lipkovich, I (2017). Multiplicity considerations in subgroup analysis. *Stat Med.* 36(28):4446-4454.
2. Guo, X and He, X (2020). Inference on selected subgroups in clinical trials, *JASA*, 116, 535; 1498-1506.
3. Freidlin, B, Sun, Z, Gray, R, Korn, E (2013). Phase III clinical trials that integrate treatment and biomarker evaluation. *JCO*
4. Ivanova, A, Israel, E, LaVange, L, Peters, M, Denlinger, LC, Moore, WC, Bacharier, LB, Marquis, MA, Gotman, NM, Kosorok, MR, Tomlinson, C, Mauger, DT, Georas, SN, Wright, RJ, Noel, P, Rosner, GL, Akuthota, P, Billheimer, D, Bleecker, ER, Cardet, JC, Castro, M, DiMango, EA, Erzurum, SC, Fahy, JV, Fajt, ML, Gaston, BM, Holguin, F, Jain, S, Kenyon, NJ, Krishnan, JA, Kraft, M, Kumar, R, Liu, MC, Ly, NP, Moy, JN, Phipatanakul, W, Ross, K, Smith, LJ, Szeffler, SJ, Teague, WG, Wechsler, ME, Wenzel, SE, White, SR (2020). The Precision Intervention in Severe and/or Exacerbation Prone Asthma (PrecISE) adaptive platform trial: statistical considerations. *Journal of Biopharmaceutical Statistics* 30(6) 1026-1037.
5. Joshi, N, Fine, J, Chu, R, Ivanova, A (2019). Estimating the subgroup and testing for treatment effect in a post-hoc analysis of a clinical trial with a biomarker. *Journal of Biopharmaceutical Statistics* 29(4), 685-695.
6. Parashar, D, Bowden, J, Starr, C, Wernisch, L, Mander, A (2016). An optimal stratified simon two-stage design. *Pharmaceutical Statistics.* 15(4):333-340.
7. Rothmann, MD, Zhang, JJ, Lu, L, Fleming, TR. (2012). Testing in a prespecified subgroup and the intent-to-treat population. *Drug Inf J.* 46(2):175-179.
8. Simon, R (2008). Development and validation of biomarker classifiers for treatment selection. *J. Statist. Planng Inf.* 138, 308–320.
9. Zhang, Z, Li, M, Lin, M, Soon, G, Greene, T, Shen, C (2017). Subgroup selection in adaptive signature designs of confirmatory clinical trials. *Journal of the Royal Statistical Society: Series C.* 66: 345-361.