Interim Analysis

Michael Martens, PhD

November 10, 2022

Stopping a Clinical Trial Early

- Why stop a trial early?
 - Strong evidence of benefit or futility of novel treatment
 - Safety concerns found through toxicity monitoring
 - Logistical issues: Interruption of drug supply, lack of funding
 - Accrual problems
- Stopping early for established benefit or for futility needs to be built into the study design to support trial's validity
- One common approach is group sequential design: efficacy compared between treatments after each group of patients are enrolled

Early Treatment Comparisons

- How do we judge whether an early treatment difference is sufficiently large or small to justify early termination?
- General approach:
 - Primary research hypothesis is specified, e.g. null hypothesis of no treatment difference
 - Efficacy data are collected over time
 - Primary outcome assessed at interim analyses
 - At each interim analysis, a test statistic is constructed and compared to a stopping boundary
 - Stopping criteria (boundaries) are chosen to preserve operating characteristics of trial (type I error rate, power)

Early Treatment Comparisons

- This general structure applies to
 - *t*-tests comparing mean between two treatments
 - Z tests comparing two proportions
 - Log-rank tests for comparing two survival curves
 - Common regression models: ANCOVA/linear regression, logistic, generalized linear, Cox proportional hazards
 - Other likelihood-based tests
- Reference: Jennison and Turnbull, Group Sequential Methods with Applications to Clinical Trials

Comparing Two Means of Normal Outcomes

- Assume we want to compare a normal outcome between two treatment groups
- Let X_{ij} denote the outcome for patient j in treatment group i
- Assume $X_{ij} \sim N(\mu_i, \sigma^2)$
- Treatment effect: $\delta = \mu_1 \mu_2$
- Null hypothesis: $H_0: \delta = 0$
- Sample size in treatment group i at time t: n_{it}
- Sample mean in treatment group *i* at time *t*:

$$ar{X}_{it} = \sum_{j=1}^{n_{it}} X_{ij}/n_{it}$$

Comparing Two Means

Test statistic:

$$Z(t) = rac{ar{X}_{1t} - ar{X}_{2t}}{\sqrt{\sigma^2/n_{1t} + \sigma^2/n_{2t}}}$$

• For large sample sizes, $Z(t) \sim N(\eta_t(\delta), 1)$, where

$$\eta_t(\delta) = \frac{\delta}{\sqrt{\sigma^2/n_{1t} + \sigma^2/n_{2t}}}$$

- Under H_0 : $\delta = 0$, $Z(t) \sim N(0,1)$
- Decision rule: Reject H_0 if $|Z(t)| \ge b(t)$, where b(t) is some critical value / boundary value which depends on t

Fixed Sample vs. Group Sequential Design

Fixed Sample Design:

- Collect observations from all patients
- Analyze results once using all observations
- Critical value of $z_{1-\alpha/2}$ used to produce type I error rate of α

Group Sequential Design:

- Collect K > 1 groups of observations during trial duration
- After each new group of observations is obtained, conduct analysis and stop trial if stopping criteria are reached
- Final sample size is a random variable; depends on when the trial stops

Group Sequential Design

- Reject H_0 the first time the boundary is crossed, i.e. first j where $|Z(t_j)| \ge b(t_j)$
- Using this strategy, we reject H_0 if

$$\{|Z(t_1)| \geq b(t_1)\} \text{ or } \{|Z(t_2)| \geq b(t_2)\} \text{ or } \cdots \text{ or } \{|Z(t_K)| \geq b(t_K)\}$$

This is equivalent to the event

$$\cup_{j=1}^K \{|Z(t_j)| \geq b(t_j)\}$$

Alternatively, we accept the null hypothesis whenever

$$\{|Z(t_1)| < b(t_1), \cdots, |Z(t_K)| < b(t_K)\}, \text{ or } \cap_{j=1}^K \{|Z(t_j)| < b(t_j)\}$$

• How to choose the boundary values $b(t_i)$?



Naive Analysis

- Suppose we reject H_0 if $|Z(t_j)| \ge z_{1-\alpha/2}$ at any interim analysis, i.e. use rejection boundaries from fixed sample design
- If $\alpha =$ 0.05, the overall type I error rate would be

$$P_{H_0}(\text{reject } H_0) = P_{H_0}\{\cup_{j=1}^K [|Z(t_j)| \ge 1.96]\} > 0.05 \text{ for } K \ge 2$$

- Multiple testing issue is at play
- Effect of multiple (equally spaced) looks on type I error rate:

K	1	2	3	4	5	10	20
α	0.050	0.083	0.107	0.126	0.142	0.193	0.246

Boundary Selection for Proper Analysis

- Need to choose boundary values carefully to protect overall type I error rate
- Must satisfy $Pig(\cup_{j=1}^K \left\{ |Z(t_j)| \geq b(t_j) \right\} ig) \leq \alpha$
- Two common boundary choices for equally spaced interim analyses (presented on Z scale):
 - Pocock Design
 - O'Brien-Fleming Design

Pocock Design

- Same boundary on Z statistic at each interim analysis: $b(t_1) = \cdots = b(t_K) = C_P(K, \alpha)$
- Pocock Design Decision Rule:
 - After group $j=1,\cdots,K-1$, stop early and reject H_0 if

$$|Z(t_j)| \geq C_P(K, \alpha);$$

- else, enroll the next cohort of patients
- After group K, reject H_0 if $|Z(t_K)| \ge C_P(K, \alpha)$; else, accept H_0
- Note that critical value C_P is higher than $z_{1-\alpha/2}$ and depends on K (increases as K does)

O'Brien-Fleming Design

- O'Brien-Fleming Decision Rule:
 - After group $j=1,\cdots,K-1$, stop early and reject H_0 if

$$|Z(t_j)| \geq C_B(K,\alpha)\sqrt{K/j};$$

else, enroll the next cohort of patients

- After group K, reject H_0 if $|Z(t_K)| \ge C_B(K, \alpha)$; else, accept H_0
- Note that critical value C_B is higher than $z_{\alpha/2}$ and depends on K (increases as K does)
- Critical values are higher at early analyses, lower at later interim analyses

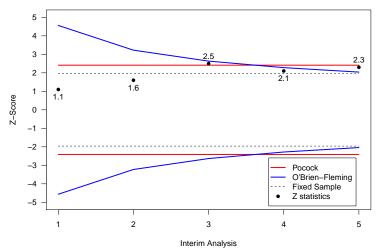
Boundaries for Pocock and O'Brien-Fleming Designs

- Values for C_P and C_B can be found in Tables 2.1 and 2.3 of Jennison and Turnbull, Table 14.2 of Piantadosi
- Software can compute these for any specified α and K values:
 - SAS PROC SEQDESIGN
 - R function gsDesign in the "gsDesign" package

 Group sequential analysis with 5 interim looks at the data, 5% type I error rate

Look	Pocock	O'Brien-Fleming		
1	2.413	$2.040\sqrt{5/1} = 4.562$		
2	2.413	$2.040\sqrt{5/2} = 3.226$		
3	2.413	$2.040\sqrt{5/3} = 2.634$		
4	2.413	$2.040\sqrt{5/4} = 2.281$		
5	2.413	2.040		

Example: Stopping boundaries with 5 interim looks, $\alpha=5\%$



Sample Size Considerations

- Group sequential trials require larger maximum sample sizes than fixed sample-designed trials
- Max sample size for group sequential design, like a fixed sample design, is proportional to $(z_{1-\alpha/2}+z_{1-\beta})^2\sigma^2/\delta^2$
- Max sample size can be written in terms of a sample size inflation factor R and fixed sample design size n_F:

$$n_{GS} = R \cdot n_F$$

- Inflation Factors depend on
 - a
 - β (equivalently, power)
 - Stopping boundary shape (e.g. Pocock, O'Brien-Fleming)
 - Number of planned looks, K

Sample Size Considerations

- Inflation factors for Pocock and O'Brien-Fleming designs can be found in Tables 2.2 and 2.4 of Jennison and Turnbull
- Software can compute these for specified α , β , K values and boundary shapes:
 - SAS PROC SEQDESIGN
 - R function gsDesign in the "gsDesign" package

- Suppose we want to test H_0 : $\mu_1 = \mu_2$ with type I error $\alpha = 0.05$ and power $1 \beta = 0.9$ to detect $\delta = \mu_1 \mu_2 = 1$, when the observations have a variance $\sigma^2 = 4$
- Fixed sample size:

$$n_F = \frac{2(z_{1-\alpha/2} + z_{1-\beta})^2 \sigma^2}{\delta^2} = \frac{2(1.96 + 1.28)^2 4}{1} = 83.98$$

$$\approx 84 \text{ patients per arm}$$

- Pocock design with 5 looks (equally spaced):
 - Inflation factor: $R_P(5, 0.05, 0.1) = 1.207$
 - Sample size needed:

$$n_{GS} = 84 \cdot 1.207 = 101.5 \approx 102 \text{ per arm}$$

- Group size to enroll for each stage is $m=102/5=20.4\approx 21$ per treatment
- Stop and reject H_0 at analysis j if $|Z(t_j)| \ge 2.413$ for $j = 1, \ldots, 5$

- O'Brien-Fleming design with 5 looks (equally spaced)
 - Inflation factor: $R_{OBF}(5, 0.05, 0.1) = 1.026$
 - Sample size needed:

$$n_{GS} = 84 \cdot 1.026 = 86.2 \approx 87 \text{ per arm}$$

- Group size to enroll for each stage is $m=87/5=17.4\approx 18$ per treatment
- Stop and reject H_0 at analysis j if $|Z(t_j)| \ge 2.040\sqrt{5/j}$ for $j=1,\ldots,5$

Expected Sample Size

- Let V be the interim analysis number at the time that the trial stops
- Let N be the trial's final sample size; note that N depends on
- Probability of stopping at analysis j is P(V = j), given by

$$P[|Z(t_1)| < b(t_1), \ldots, |Z(t_{j-1})| < b(t_{j-1}), |Z(t_j)| \ge b(t_j)]$$

- Let E(V) be the expected number of interim analyses conducted assuming H_1 is true $(\delta \neq 0)$
- Expected sample size if H_1 is true:

$$E(N) = n_F \cdot R \cdot \frac{E(V)}{K}$$



Expected Sample Size

• For a design with $\alpha=$ 0.05, $1-\beta=$ 0.9, and K= 5:

Boundary	E(V)	Max N	E(N)
Pocock	2.83	$R_P \cdot n_F = 1.21 n_F$	0.68 <i>n_F</i>
OBF	3.65	$R_{OBF} \cdot n_F = 1.03 n_F$	0.75 <i>n_F</i>
Fixed		n_F	n_F

• O'Brien-Fleming designs have a lower maximum sample size than Pocock, but higher average sample size if H_1 is true

Choosing a Stopping Boundary

- Pocock Design:
 - Higher chance of early stopping under alternative
 - Higher inflation factor and maximum N
 - Lower expected sample size under alternative
 - Higher expected sample size under H_0
- O'Brien-Fleming Design:
 - Lower chance of stopping early under alternative
 - Lower inflation factor and maximum N
 - Higher expected sample size under alternative
 - Lower expected sample size under H₀

Choosing a Stopping Boundary

- O'Brien-Fleming (OBF) design is much more common than Pocock; reasons include:
 - Many trials fail to find efficacy, in which case N is smaller and trial is less costly with OBF design
 - In early stages of trial, data may be less reliable and possibly un-representative; better to make it more difficult to stop early
 - Similarly, may prefer to make it easier to reject later in the trial when more data are available
 - OBF still offers big reduction in expected sample size under H₁ with very slight increase in maximum N compared to fixed sample design

Choosing a Stopping Boundary

- For OBF test, boundary value at the end of the study is very close to usual fixed sample boundary
- Example:
 - Suppose Z = 2.30 at end of study
 - For fixed sample design, $|Z| \ge 1.96$, so reject H_0
 - For Pocock design with 5 looks, $|Z| \ge C_P = 2.413$, so accept H_0
 - For OBF design with 5 looks, $|Z| \ge C_B = 2.04$, so reject H_0 ;
 - OBF final boundary of 2.04 is very close to fixed sample design boundary of 1.96

Theory of Boundary Selection

Boundary values must satisfy

$$P_{H_0}[|Z(t_1)| < b(t_1), \ldots, |Z(t_K)| < b(t_K)] \ge 1 - \alpha$$

- Need to known the joint distribution of $(Z(t_1), \ldots, Z(t_K))$ to pick boundaries
- Recall that the variance of $ar{X}_{1j} ar{X}_{2j}$ is

$$V_j = \sigma^2/n_{1j} + \sigma^2/n_{2j}$$

where n_{ij} is the sample size in group i at time t_j

• Define the information at t_j as $I_j = V_j^{-1}$; then

$$Z_j = Z(t_j) = (\bar{X}_{1j} - \bar{X}_{2j})\sqrt{I_j}$$



Theory of Boundary Selection

- The vector $\mathbf{Z} = (Z_1, \dots, Z_K)$ is multivariate normally distributed
- It can be shown that:
 - $Z_j \sim N(\delta \sqrt{I_j}, 1)$ for j = 1, ..., K
 - $Cor(Z_i, Z_j) = Cov(Z_i, Z_j) = \sqrt{I_i/I_j}$ for $i, j = 1, ..., K, i \le j$, since $Var(Z_j) = 1$ for all j
- This multivariate normal structure is sometimes referred to as the Canonical Distribution; also called the Independent Increments property

Theory of Boundary Selection

- Evaluation of type I error rate requires integration of the multivariate normal distribution over subsets of \mathbb{R}^K
- This can be done efficiently by numerical methods if the independent increments property holds
- Test statistics for many common tests approximately follow the canonical distribution, so boundaries can be computed by standard methods

Boundary Selection for Other Test Statistics

 Comparison of two proportions: Z test at interim look j has test statistic

$$Z=(\hat{
ho}_{1j}-\hat{
ho}_{2j})\sqrt{I_j}, ext{ where} \ I_j=[ar{
ho}_j(1-ar{
ho}_j)(1/n_{1j}+1/n_{2j})]^{-1}$$
 ;

information is approximately proportional to sample size

- Comparison of two survival curves: Log-rank test
 - Information is approximately proportional to the number of deaths

Timing of Interim Analyses

- O'Brien-Fleming and Pocock designs assume that interim analyses occur after equal amounts of information (or sample sizes) are collected
- It may be impractical to monitor the data at equal increments of information:
 - Enrollment pattern may be unpredictable
 - Scheduling interim analyses on a calendar basis (e.g. yearly) is often more convienent; likely won't align with equal information increments
 - Nuisance parameters may be misspecified (e.g. we assumed $\sigma^2=4$, but really $\sigma^2=9$)
- The error rate α can be controlled while allowing flexibility in analyses timings using an **Error Spending Function**

Error Spending Approach for Group Sequential Trials

		Stopping	Cumulative
Time	Rejection Region	Prob (H_0)	Stopping Prob (H_0)
t_1	$\{ Z(t_1) \geq b_1\}$	θ_1	$\alpha_1 = \theta_1$
t_2	$\{ Z(t_1) < b_1, Z(t_2) \geq b_2\}$	θ_2	$\alpha_2 = \theta_1 + \theta_2$
:	:	:	:
t_{K}	$\{ Z(t_1) < b_1, \ldots,$	θ_{K}	$\alpha_{\kappa} = \theta_1 + \ldots + \theta_{\kappa}$
	$ Z(t_{K-1}) < b_{K-1}, Z(t_K) \ge b_K$	~ ^ ^	$\begin{bmatrix} \alpha_{K} - \sigma_1 + \dots + \sigma_{K} \end{bmatrix}$

- θ_j is the probability of stopping study at analysis j under H_0
- α_j is the probability of stopping study at or before analysis j under H_0
- α_j is the amount of type I error "spent" through time t_j
- Set overall significance level $\alpha_K = \alpha$

Error Spending Approach

- Lan & DeMets suggested parametrizing group sequential tests through α_j rather than boundaries b_j
- ullet A 1-1 correspondence exists between the b_j and $lpha_j$
- To determine the α_j , specify an error spending function $\alpha(\tau)$, an increasing function with $\alpha(0) = 0$, $\alpha(1) = \alpha$
- $\alpha(\tau)$ is associated with an information time scale, τ
- $\tau_j = I(t_j)/I(t_K)$ is the information fraction at analysis j
- Set $\alpha_j = \alpha(\tau_j)$ for *j*th interim analysis
- $\tau_K = 1$ represents the final analysis, with $\alpha(\tau_k) = \alpha$
- Error spending approach is equivalent to specifying a group sequential boundary, while allowing flexibility in analysis timing



Commonly Used Error Spending Functions

"Pocock type" spending function (Lan & DeMets):

$$\alpha(t) = \min \left(\alpha \log(1 + (e - 1)t), \alpha \right)$$

"O'Brien-Fleming type" spending function (Lan & DeMets):

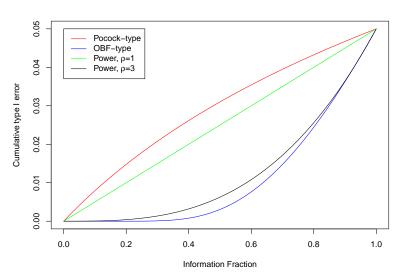
$$\alpha(t) = \min\left(4 - 4\Phi(z_{\alpha/4}/\sqrt{t}), \alpha\right)$$

Power family (Jennison & Turnbull):

$$\alpha(t) = \min(\alpha t^{\rho}, \alpha)$$
, where $\rho > 0$

ho=1 is similar to Pocock, ho=3 similar to OBF

Commonly Used Error Spending Functions



Calculation of Boundaries

- Suppose analyses will be conducted at times (t_1, \ldots, t_K)
- K and times need not be specified in advance
- Boundary values are computed iteratively according to

$$P_{H_0}\{|Z(t_1)| \ge b_1\} = \theta_1 = \alpha(\tau_1),$$

$$P_{H_0}\{|Z(t_1)| < b_1, |Z(t_2)| \ge b_2\} = \theta_2 = \alpha(\tau_2) - \alpha(\tau_1),$$

$$\vdots$$

$$P_{H_0}\{|Z(t_1)| < b_1, \ldots, |Z(t_{K-1})| < b_{K-1}, |Z(t_K)| \ge b_K\} = \theta_K = \alpha(\tau_K) - \alpha(\tau_{K-1})$$

• Can solve these recursively for b_1 , then b_2 , etc. using the multivariate normal distribution of **Z**



Considerations for Error Spending Approach

- Type I error rate is not controlled if the choice of monitoring times is dependent on the observed treatment differences
 - Interim analysis times must be chosen independent of treatment effect info
- Boundaries cannot be computed in advance unless information fractions are prespecified
- ullet For study design, we need to provide adequate power to detect δ of interest
 - Power depends on the information sequence $\{\tau_j\}$
 - Protocol should specify an information sequence to be targeted for the study (e.g. 3 looks at 50%, 75%, 100% information)
 - An equally spaced sequence is often assumed when designing study

Impact of Misspecification / Change of Design Parameters

- If design parameters are misspecified or changed, the power can be impacted:
 - Number of analyses K: minimal impact
 - Information fractions: minimal, if close to the sequence assumed at the design stage
 - Nuisance parameters: can result in over or underpowered study; same problem happens with fixed sample designs

Information Fractions

- Two potential definitions of information fraction:
 - Ratio of sample size at interim analysis to maximum sample size
 - Ratio of information at interim analysis to maximum information, where information = $Var(trt \text{ effect estimate})^{-1}$

Example

- Clinical trial to test H_0 : $\mu_1 = \mu_2$ with $\alpha = 0.05$, power $1 \beta = 0.9$ when $\mu_1 \mu_2 = 1$, and variance $\sigma^2 = 4$
- Fixed sample design requires n = 84 per group
- Monitor 3 times, equal information, $\rho=3$ spending function, $\alpha(\tau)=\min(\alpha\tau^3,\alpha)$: Inflation factor $R_{\rho=3}=1.018$
- Maximum sample size: $n_{
 ho=3}=84\cdot 1.018=85.5pprox 86$ per group
- Maximum Information = SE^{-2} :

$$MI = \left[\sigma^2(1/n_1 + 1/n_2)\right]^{-1} = \frac{86/2}{4} = 10.8$$



Group	n	x	S
1	20	7.1	2.5
2	25	8.3	2.9

- Strategy A: Information fraction based on ratio of sample sizes
 - Information fraction: $\tau_1 = 45/172 = 0.26$
 - Type I error spent: $\alpha(\tau_1) = 0.05(0.26)^3 = 0.00088$
 - Boundary: $b_1 = 3.326$
- Test statistic:

$$Z(t_1) = \frac{7.1 - 8.3}{0.8055} = -1.490$$

Continue study to next analysis

Group	n	\bar{x}	S
1	20	7.1	2.5
2	25	8.3	2.9

- Strategy B: Information fraction based on ratio of current information to maximum information
 - Standard Error: $SE = \sqrt{s_1^2/n_1 + s_2^2/n_2} = 0.8055$
 - Information: $I(t_1) = (0.8055)^{-2} = 1.541$
 - Information fraction: $\tau_1 = 1.541/10.8 = 0.143$
 - Type I error spent: $\alpha(\tau_1) = 0.05(0.143)^3 = 0.00015$
 - Boundary: $b_1 = 3.791$
- Test statistic:

$$Z(t_1) = -1.490$$

Continue study to next analysis



Group	n	\bar{x}	S
1	55	6.9	2.8
2	60	8.4	3.1

- Strategy A: Information fraction based on ratio of sample sizes
 - Information fraction: $\tau_2 = 115/172 = 0.669$
 - Cumulative Type I error spent: $\alpha(\tau_2) = 0.05(0.669)^3 = 0.015$
 - Type I error spent at this look: $\theta_2=0.015-0.00088=0.0141$
 - Boundary: $b_2 = 2.452$
- Test statistic:

$$Z(t_2) = \frac{6.9 - 8.4}{0.5502} = -2.726$$

Stop study and reject H₀



Group	n	\bar{x}	S
1	55	6.9	2.8
2	60	8.4	3.1

- Strategy B: Information fraction based on ratio of current information to maximum information
 - Standard Error: $SE = \sqrt{s_1^2/n_1 + s_2^2/n_2} = 0.5502$
 - Information: $I(t_2) = (0.5502)^{-2} = 3.303$
 - Information fraction: $\tau_2 = 3.303/10.8 = 0.306$
 - Cumulative Type I error spent: $\alpha(\tau_2) = 0.05(0.306)^3 = 0.0014$
 - Type I error spent at this look:
 - $\theta_2 = 0.0014 0.00015 = 0.00125$
 - Boundary: $b_2 = 3.227$
- Test statistic: $Z(t_2) = -2.726$
- Continue study to next analysis



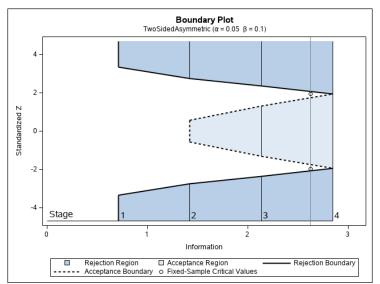
Example

- Some evidence exists that the prespecified variance is wrong: assumed $\sigma^2 = 4$ but $s_1^2 = 7.84, s_2^2 = 9.61$ at look 2
- With $\sigma_1^2=7.84$ and $\sigma_2^2=9.61$ we would need 186 patients per arm to meet the power requirement
- Strategy B allows for sample size escalation to attain the desired power, since very little type I error is spent early
- Strategy A spends the type I error based on the original assumption about the variance, so will be underpowered

Group Sequential Design: Extensions

- Early futility stopping:
 - Futility boundaries $\{d_j\}$ included such that trial stops for futility at stage j if $|Z(t_j)| \leq d_j$
 - With equal information increments, these can be determined directly similar to Pocock and O'Brien-Fleming designs
 - Error spending approach: Specify function $\beta(\tau)$ such that $\beta(0) = 0$, $\beta(1) = \beta$; use it to determine futility boundaries at interim analyses (type II error spending)
- One-sided testing
- Noninferiority and equivalence testing

Example: Group Sequential Design with Early Efficacy and Futility Stopping



Group Sequential Design: Extensions

- Other types of outcomes (binary, categorical, survival, etc.): Same boundary specification and error-spending methods can be used, relying on asymptotic normal distributions of treatment effects for most common methods, including:
 - Pearson's chi-square and Mantel-Haenszel tests
 - Logistic regression model
 - Log rank test and Cox proportional hazards models
 - Generalized linear models
 - Linear and generalized liear mixed models

Stochastic Curtailment

- Idea: Terminate a study if a particular outcome is highly likely given the observed data
- Often this is done using a concept called conditional power
- Testing $H_0: \theta = 0$ vs. $H_1: \theta = \delta$, where θ represents the treatment effect
- Reference test $\mathcal T$ used with type I error α under $\theta=0$ and power $1-\beta$ at $\theta=\delta$
 - One-sided, fixed-sample test
 - Reject H_0 if $Z \geq z_{\alpha}$
 - Power of $1-\beta$ achieved by choosing the sample size to yield an information level I_K

Stochastic Curtailment

- At interim analysis j, let Z_j denote the test statistic so far
- Conditional Power to detect θ at stage j:

$$p_j(\theta) = P_{\theta}(\mathcal{T} \text{ rejects } H_0 | \text{data at stage } j) = P_{\theta}(\mathcal{T} \text{ rejects } H_0 | Z_j)$$

- If $p_j(0)$ is high, then the reference test is likely to reject H_0 even if H_0 is true
- If $p_j(\delta)$ is low, then the reference test is unlikely to reject H_0 even if H_0 is false
- Formal monitoring rules:
 - Early termination for efficacy: Reject H_0 at stage j if $p_j(0) \ge \gamma$ (Usually 0.8 or 0.9)
 - Early termination for futility: Accept H_0 at stage j if $1 p_j(\delta) \ge \gamma'$



Stochastic Curtailment: Properties

- Fully sequential monitoring is permitted (can look anytime)
- Can be shown that the type I error rate is no more than α/γ , type II error rate is no more than β/γ'
- Can ensure appropriate α,β error rates for sequential procedure by choosing fixed sample test $\mathcal T$ to have type I error rate $\alpha\gamma$ and power $1-\beta\gamma'$ at $\theta=\delta$

Conditional Power

• Conditional power at interim analysis j, with information l_j

$$p_j(\theta) = P_{\theta}(Z_K \ge z_{1-\alpha}|Z_j)$$

• The conditional distribution of $Z_K|Z_j$ is

$$Z_K|Z_j \sim N\left(Z_j\sqrt{I_j/I_K} + rac{\theta(I_K - I_j)}{\sqrt{I_K}}, 1 - rac{I_j}{I_K}
ight)$$

Then the conditional power is

$$p_j(\theta) = \Phi\left(\frac{Z_j\sqrt{I_j} - z_{1-\alpha}\sqrt{I_K} + \theta(I_K - I_j)}{\sqrt{I_K - I_j}}\right)$$

Conditional Power

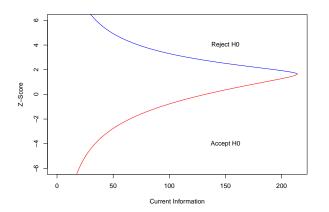
- Stopping rules can be directly translated into stopping boundaries:
 - Stop and reject H_0 if $p_j(0) \ge \gamma$

$$Z_j \ge z_{1-\alpha} \sqrt{I_K/I_j} + z_\gamma \sqrt{(I_K - I_j)/I_j}$$

• Stop and accept H_0 if $p_j(\delta) \leq 1 - \gamma'$

$$Z_j \leq z_{1-\alpha} \sqrt{I_K/I_j} + z_{1-\gamma'} \sqrt{(I_K - I_j)/I_j} - \frac{\delta(I_K - I_j)}{\sqrt{I_j}}$$

Stochastic Curtailment Example: $\alpha = 0.05$, $1 - \beta = 0.9$, $\delta = 0.2$, $I_K = 214.1$, $\gamma = \gamma' = 0.8$



- Actual type I error rate $\leq \alpha/\gamma = 0.0625$
- Actual power is $\geq 1 0.1/0.8 = 0.875$



Two-sided Conditional Power

- Reference test: Reject H_0 if $|Z_K| \ge z_{\alpha/2}$
- Conditional power at analysis j:

$$p_{j}(\theta) = \Phi\left(\frac{Z_{j}\sqrt{I_{j}} - z_{\alpha/2}\sqrt{I_{K}} + \theta(I_{K} - I_{j})}{\sqrt{I_{K} - I_{j}}}\right) + \Phi\left(\frac{-Z_{j}\sqrt{I_{j}} - z_{\alpha/2}\sqrt{I_{K}} - \theta(I_{K} - I_{j})}{\sqrt{I_{K} - I_{j}}}\right)$$

 Can use this for fully sequential testing, with same impacts on type I/II error rates

Comments on Conditional Power

- Offers a simple rule that is often applied for early futility stopping
- Can be compared directly to group sequential methods as long as we calibrate the error rates to be equal
- Fully sequential, not group sequential monitoring
 - Allows for arbitrary, unplanned analysis times, and any K
 - May be conservative (α too small) compared to a group sequential design with a planned analysis schedule

- Multiple looks at the data affect confidence intervals, just like they affect significance levels of hypothesis tests
- Unadjusted (naive) 95% confidence intervals for mean differences: After each group j of observations, $j=1,\ldots,K$, form a 95% confidence interval

$$ar{X}_{1j} - ar{X}_{2j} \pm 1.96 \sqrt{\sigma_1^2/n_{1j} + \sigma_2^2/n_{2j}}$$

• The probability that all K intervals contain the true value of θ is <95%

• Repeated Confidence Intervals: Set of random intervals $\{RCI_j\}$ where the simultaneous coverage probability is maintained at $1-\alpha$:

$$P(\theta \in \mathsf{RCI}_j \text{ for all } j = 1, \dots, K) \ge 1 - \alpha$$

- RCIs constructed by inverting group sequential tests
- Define $Z_j(\theta_0) = Z_j \theta_0 \sqrt{I_j}$ for j = 1, ..., K
- Group sequential test: reject $H_0: \theta = \theta_0$ at stage j if $|Z_j(\theta_0)| \ge c_j(\alpha)$
- Then the RCIs are defined as

$$RCI_{j} = \{\theta_{0} : |Z_{j}(\theta_{0})| < c_{j}(\alpha)\} = \left(\frac{Z_{j} - c_{j}(\alpha)}{\sqrt{I_{j}}}, \frac{Z_{j} + c_{j}(\alpha)}{\sqrt{I_{j}}}\right)$$
$$= \left(\hat{\theta}_{j} - c_{j}(\alpha)I_{j}^{-1/2}, \hat{\theta}_{j} + c_{j}(\alpha)I_{j}^{-1/2}\right),$$

where $\hat{ heta}_j$ is the estimate of heta at look j



- For comparing two means:
 - Naive 95% confidence interval:

$$\bar{X}_{1j} - \bar{X}_{2j} \pm z_{1-\alpha/2} \sqrt{\sigma_1^2/n_{1j} + \sigma_2^2/n_{2j}}$$

RCIs:

$$\bar{X}_{1j} - \bar{X}_{2j} \pm c_j(\alpha) \sqrt{\sigma_1^2/n_{1j} + \sigma_2^2/n_{2j}}, \quad j = 1, ..., K$$

 With O'Brien-Fleming design, RCIs are wide at early analyses and get closer to naive CIs at end of study

Example: Group Sequential Design in R

- Designing a group sequential study:
 - H_0 : $\mu_1 = \mu_2$, 2-sided $\alpha = 0.05$, power $1 \beta = 0.9$ when $\delta = \mu_1 \mu_2 = 1$, $\sigma^2 = 4$
 - K = 3 looks with an O'Brien-Fleming boundary
 - Sample size for fixed design

$$n = \frac{2(z_{1-\alpha/2} + z_{1-\beta})^2 \sigma^2}{\delta^2} = 84$$
 per group, 168 total

• Sample size for O'Brien-Fleming design using R:

gsDesign(k=3, test.type=2, sfu="OF",n.fix=168)

Example: O'Brien-Fleming Design in R

Symmetric two-sided group sequential design with 90 % power and 2.5 % Type I Error. Spending computations assume trial stops if a bound is crossed.

```
Analysis N Z Nominal p Spend

1 57 3.47 0.0003 0.0003

2 114 2.45 0.0071 0.0069

3 171 2.00 0.0225 0.0178

Total 0.0250
```

Example: O'Brien-Fleming Design in R

```
Boundary crossing probabilities and expected sample size
assume any cross stops the trial
Upper boundary (power or Type I Error)
         Analysis
  Theta 1 2 3 Total E{N}
 0.0000 0.0003 0.0069 0.0178 0.025 169.9
 0.2501 0.0565 0.5288 0.3147 0.900 134.2
Lower boundary (futility or Type II Error)
         Analysis
  Theta 1 2 3 Total
 0.0000 3e-04 0.0069 0.0178 0.025
 0.2501 0e+00 0.0000 0.0000 0.000
```

++ alpha spending:

O'Brien-Fleming boundary.

Example: O'Brien-Fleming Design

• Interim analysis 1:

Group	n	x	S
1	20	7.1	2.5
2	25	8.3	2.9

- Test statistic $Z(t_1) = -1.49$
- Boundary: $b_1 = 3.47$
- Continue study

Example: O'Brien-Fleming Design

• Interim analysis 2:

Group	n	\bar{x}	S
1	55	6.9	2.8
2	60	8.4	3.1

- Test statistic: $Z(t_2) = -2.73$
- Boundary: $b_2 = 2.45$
- Stop study and reject H_0

- 95% RCIs:
 - Look 1: Boundary value $b_1(\alpha) = 3.47$

$$\mathsf{RCI}_1 = (7.1 - 8.3) \pm 3.47 (0.8055) = [-4.00, 1.60]$$

• Look 2: Boundary value $b_2(\alpha) = 2.45$

$$\mathsf{RCI}_2 = (6.9 - 8.4) \pm 2.45 \\ (0.5502) = [-2.85, -0.15]$$