

Early Phase Trial Designs and Estimand

Yuan Ji, PhD

Program for Computational Genomics & Medicine
NorthShore University HealthSystem
Department of Public Health Sciences
The University of Chicago

SBF 2017

December 4, 2017

COI Disclosure

- Consultants for Seattle Genetics and Biogen.
- Co-Founder of Laiya Consulting, Inc., a statistical solution and consulting company

ESTIMAND

An ESTIMAND includes

1. Target population
2. Endpoint (the variable to be measured)
3. Strategies to account for **intercurrent events** (potential confounders)
4. **Population summaries** for statistical inference and comparison

ESTIMAND for confirmatory (late-phase) trials depends on results from early-phase trials, such as

- target population

There could be discrepancies between ESTIMAND for confirmatory trials and important components for early-phase trials, such as

- endpoint

Phase I dose-finding (Oncology)

Consider trials with fixed doses.

Setup Climb **up and down** a sequence of D ordered doses of a new drug to determine the maximum tolerated dose (MTD).

Data At each dose i , n_i patients are tested, y_i patients experienced **toxicity outcome** (DLT).

Parameters Dose i has a toxicity probability of p_i (unknown).

Sampling Model Binomial $y_i | p_i \sim \text{Bin}(n_i, p_i)$

Assumption **Toxicity Monotonicity** : $p_i \leq p_{i+1}$.

Hidden assumption **Efficacy Monotonicity** : $q_i \leq q_{i+1}$ – if not, why escalate when the dose is safe?

Goal to find the **MTD**, defined as the highest dose with **toxicity rate** lower (or close to) **a target rate**, p_T , e.g., $p_T = 0.30$.

Intercurrent event **Death** (how to account for death in dose-finding trials)

ESTIMAND for dose-finding trials

1. Target Population

- Traditionally, all comers (e.g., all patients with solid tumors)
- Recently, in Immune-Oncology (e.g., CAR-T trials), targeted cancer types (subtypes). Often the same as late-phase population

2. Endpoint

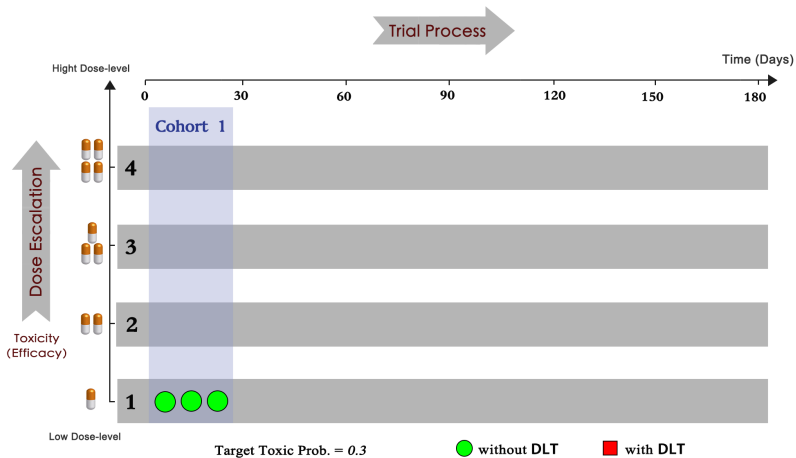
- Binary DLT: yes/no
- Different grades; multiple cycles; time-to-event

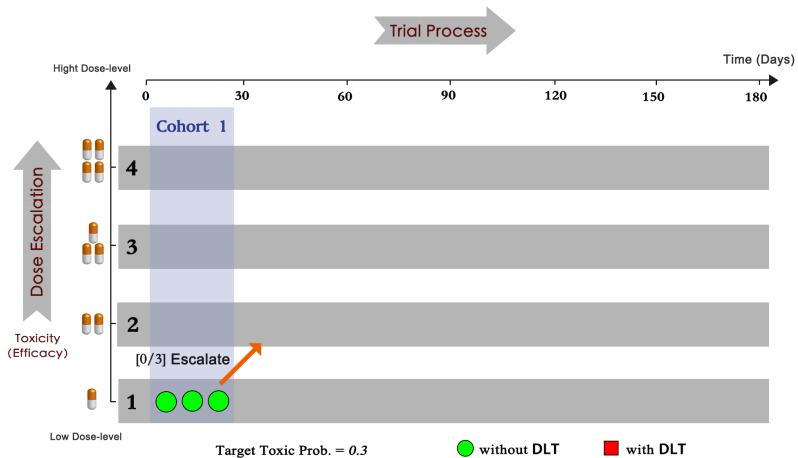
3. Intercurrent Event

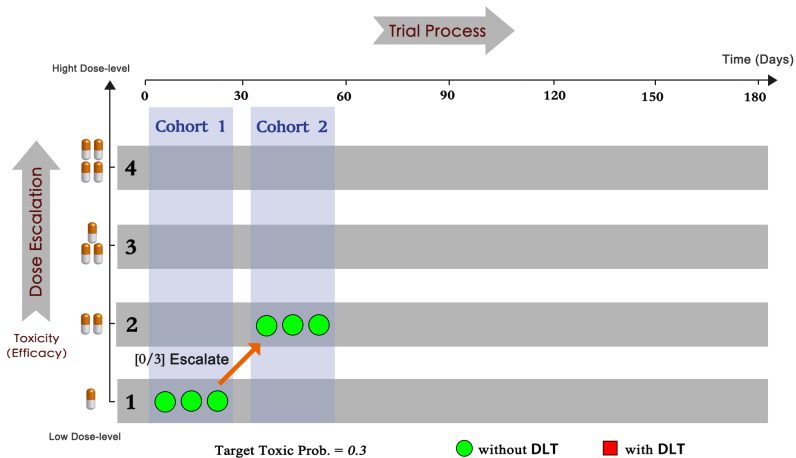
- Death: due to toxicity vs. due to disease
- Side effects from other medications

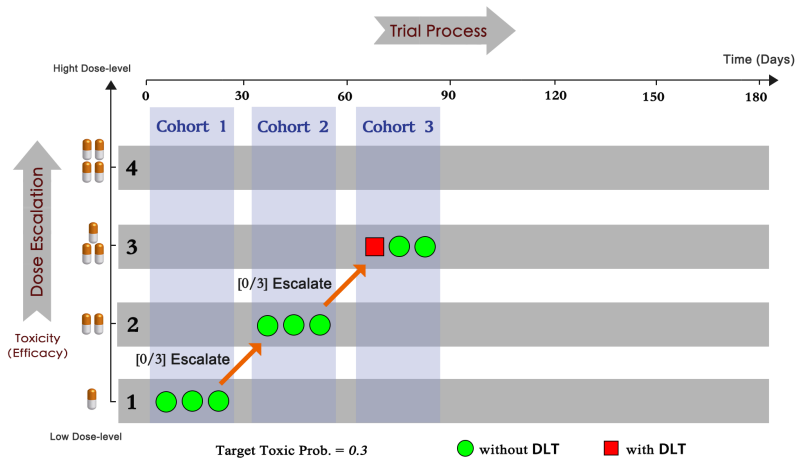
4. Population Summary

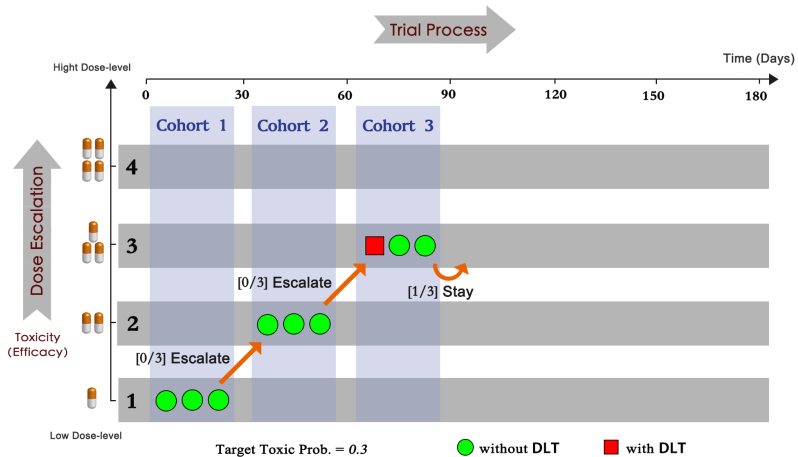
- DLT rates at different doses
- Total toxicity risk score (for toxicity grades)
- Average toxicity rates over cycles

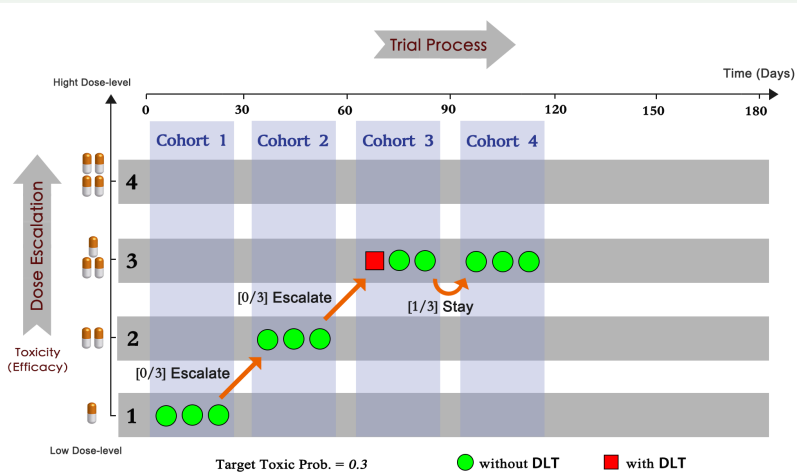


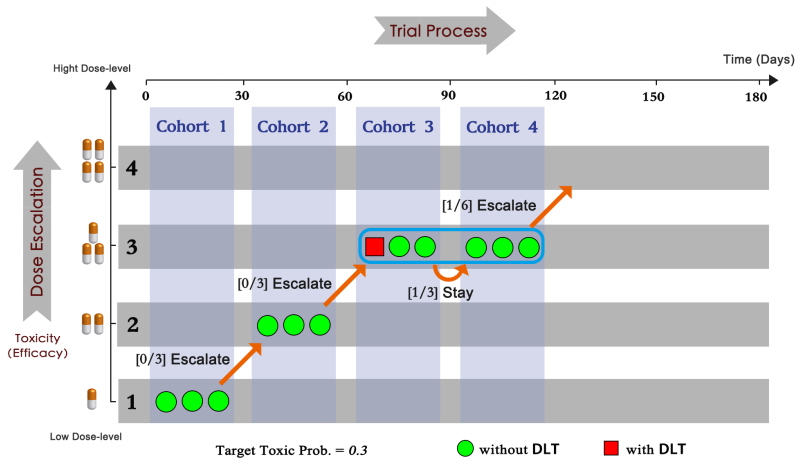


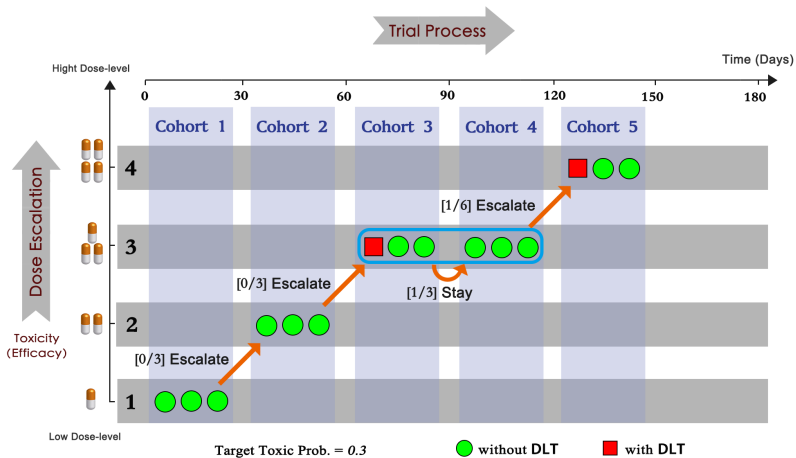


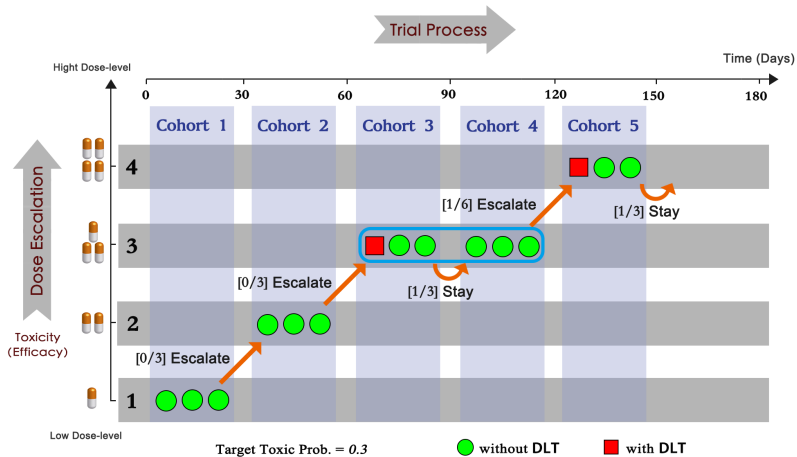


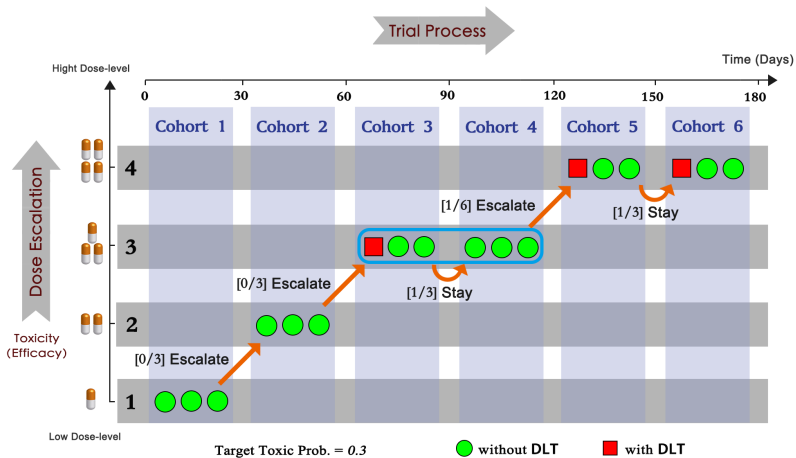


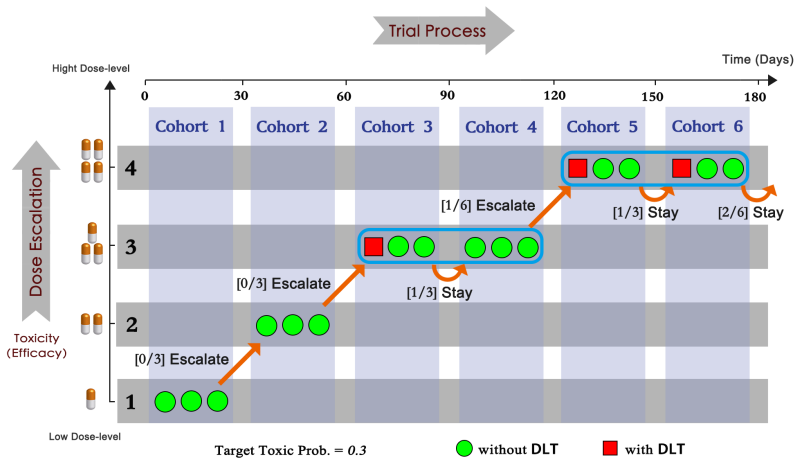








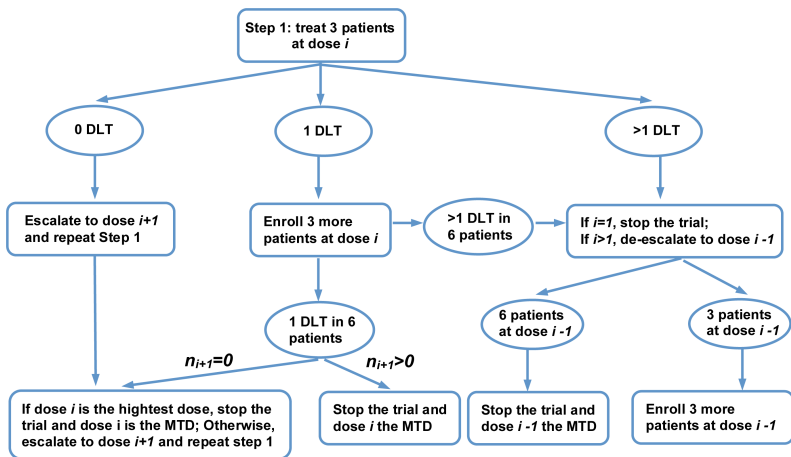




Existing Designs

- 3+3
 - Storer (1989). **Algorithmic design**; simple and transparent; popular among physicians
 - Lacks a standard program; performs worse than model-based designs
- CRM
 - The first **model-based design**. First publication in 1990.
 - Many different versions; **black-box to physicians**
- mTPI and mTPI-2
 - Ji et al. (2010); Guo et al. (2017). **Model-based interval design** in an algorithmic presentation
 - Getting popular in the community; user-friendly software; simple and transparent
- CCD and BOIN
 - CCD (Ivanova et al. 2007), extension of CCD – BOIN (Liu and Ying 2015); also **model-based interval designs** but with a different framework
 - Simple inference based on point estimate of toxicity probabilities; BOIN asymptotic behavior is strange

The Industry-Standard 3+3 Design



– Yang et al. (2015)

The 3+3 Design

Standard in clinical community (simple, transparent, “make sense”, perform reasonably well if $p_T \sim 17\%$ or 30% .)

Favorite Target to statisticians

- Not model-based
- No more than 6 patients per dose
- Arbitrary choice of “3”
- Conservative – slow escalation

Dominant in practice (e.g., Rogatko et al. 2007)

- $>98\%$ of all phase I trials between 1991-2006 were based on 3+3 or its variations
- Out of 1,235 trials during the period, 20 were based on CRM; 3 based on EWOC (a variation of CRM)
- getting less popular and more frequently criticized recently (Nie et al., 2016)

The 3+3 Design: Is it conservative?

- Ji and Wang (2013, *JCO*) showed that with matched sample size, **3+3 is less safe and reliable when compared to the mTPI design**, a model-based design.
- The 2015 FDA/AACR Dose-Finding Symposium concluded that (Nie et al., 2016, *Clinical Cancer Research*)

*“The MTD/3+3 approach is not optimal and may result in recommended doses that are **unacceptably toxic** for many patients and in **dose reduction/interruptions** that might have an impact on effectiveness.”*

The CRM Design – a specific model

Perhaps the most popular version of the CRM is the power model:

- The **dose-response curve** : $p_i = p_{i0}^{\exp(\alpha)}$, where p_{i0} are fixed and prespecified constants, and α is a parameter that describes the dose response curve.
- The prior for α is $N(0, 2)$.
- The p_{i0} 's are decided by solving $E[p_i^{\exp(\alpha)}] = s_i$, where s_i 's are a set of prior probabilities that one must determine (called "**skeletons**").
- A binomial likelihood: $\prod_{i=1}^d p_i^{y_i} (1 - p_i)^{n_i - y_i}$.
- Posterior of α is obtained by numerical integration.
- The next dose is $\arg \min_i | \hat{p}_i - p_T |$, where \hat{p}_i is the posterior mean.

The CRM Design – trial conduct

- Challenging to implement in practice (logistic and people issues)
- How does one actually conduct a practical trial using CRM?:
 - Need to set up an infrastructure between statisticians and nurses at clinics (potentially multiple sites).
 - Anti-CRM ad-hoc rules: Coherence and Over-dose control (e.g., no skipping dose when escalation)
 - Team meetings are needed for every patient allocation – CRM decisions may be overruled.

The mTPI and mTPI-2 designs are interval designs. The CCD and BOIN designs are a different type of interval designs.

Hallmark of "Interval Designs"

The decision of dose finding involves inference based on toxicity probability intervals.

- **Interval designs** : up-and-down decisions based on intervals (mTPI, mTPI-2, CCD, BOIN)

$$\left| \begin{array}{c} \text{Stay} \\ p_i \in (p_T - \epsilon_1, p_T + \epsilon_2) \end{array} \right| \left| \begin{array}{c} \text{Escalate} \\ p_i \in (0, p_T - \epsilon_1) \end{array} \right| \left| \begin{array}{c} \text{De-escalate} \\ p_i \in (p_T + \epsilon_2, 1) \end{array} \right|$$

- **Non-interval designs** : CRM chooses the dose

$$\arg \min_i |\hat{p}_i - p_T|,$$

3+3 uses up-and-down decisions based on

$$\frac{y_i}{n_i},$$

The mTPI Design

Interval designs are a simple, transparent, intuitive and **model-based**.

Sample size = 21 ; Target probability = 30% ; epsilon 1 = 0.05 ; epsilon 2 = 0.05 ; OLD

E: Escalate to the next higher dose; S: Stay at the same dose; D: De-escalate to the previous lower dose; DU: De-escalate to the previous lower dose and the current dose will never be used again in the trial;

Number of Patients

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21
0	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E
1	D	S	S	S	S	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E
2		DU	D	S	S	S	S	S	S	S	E	E	E	E	E	E	E	E	E	E	E
3			DU	DU	D	S	S	S	S	S	S	S	S	S	S	E	E	E	E	E	E
4				DU	DU	DU	D	D	S	S	S	S	S	S	S	S	S	S	S	S	E
5					DU	DU	DU	DU	DU	D	S	S	S	S	S	S	S	S	S	S	S
6						DU	DU	DU	DU	DU	DU	D	S	S	S	S	S	S	S	S	S
7							DU	DU	DU	DU	DU	DU	DU	D	S	S	S	S	S	S	S
8								DU	DU	DU	DU	DU	DU	DU	DU	DU	D	S	S	S	S
9									DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	S
10										DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	DU
11											DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	DU
12												DU	DU	DU	DU	DU	DU	DU	DU	DU	DU
13													DU	DU	DU	DU	DU	DU	DU	DU	DU
14														DU	DU	DU	DU	DU	DU	DU	DU
15															DU	DU	DU	DU	DU	DU	DU
16																DU	DU	DU	DU	DU	DU
17																	DU	DU	DU	DU	DU
18																		DU	DU	DU	DU
19																			DU	DU	DU
20																				DU	DU
21																					DU

Number of Toxicity

– An mTPI decision table

but in an **algorithmic form**

Interval-Based Decision Rules

Divid $(0, 1)$ into three intervals:

$$(0, p_T - \epsilon_1), \quad \underbrace{(p_T - \epsilon_1, p_T + \epsilon_2)}_{\text{Equivalence Interval}}, \quad (p_T + \epsilon_2, 1)$$

Associate with

E ,

S ,

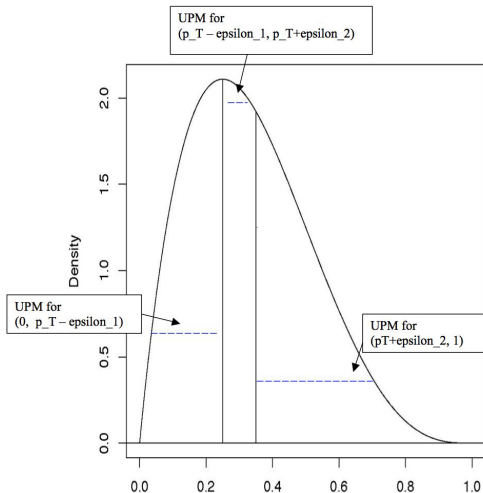
D

Measure the **unit probability mass (UPM)** of each interval under the posterior of p_i .

Decide the action corresponding to the largest UPM (Bayes rule – more on this next).

UPM and Bayes rule

$$\text{UPM (interval)} = \frac{\text{post. prob } \{ p_i \in (\text{interval}) \}}{\text{length (interval)}}$$



Ockham's razor and interval length

In mTPI, when 3 out of 6 patients have DLT and if $p_T = 0.3$, the decision based on UPM is S , to stay at the current dose. Why?

Ockham's razor states the principle that an explanation of the facts should be no more complicated than necessary

Bayesian model selection requires a prior $p(M_k)$ for the candidate model k and a prior $p(\theta_k | M_k)$. Models are selected based on $p(M_k | data)$ and automatically applies the Ockham's razor: when two models fit the data equally well, the smaller one wins.

The mTPI design considers three intervals that partition the sample space $(0, 1)$ for the probability of toxicity p_d at a given dose d :

$$\begin{aligned}
 M_E : \quad & p_d \in (0, p_T - \epsilon_1) \\
 M_S : \quad & p_d \in (p_T - \epsilon_1, p_T + \epsilon_2) \\
 M_D : \quad & p_d \in (p_T + \epsilon_2, 1)
 \end{aligned} \tag{1}$$

Typically, p_T ranges from 0.1 to 0.3 in phase I trials, and ϵ 's are usually small, say ≤ 0.05 . So M_S the middle interval is the smallest (shortest).

Ockham's razor and interval length – Con't

So mTPI is based on the decision rule for dose i

$$\mathcal{D}_{\text{mTPI}} = \arg \max_{k \in \{E, S, D\}} UPM(k, i) \quad (2)$$

Ockham's razor and interval length – Con't

So mTPI is based on the decision rule for dose i

$$\mathcal{D}_{\text{mTPI}} = \arg \max_{k \in \{E, S, D\}} UPM(k, i) \quad (2)$$

Turns out this rule is the Bayes rule for the following decision framework:

Theorem 1. Given the sampling model $y_i | p_i \sim \text{Bin}(n_i, p_i)$ and priors

$$f(p_i | M_k) \sim \frac{1}{S(M_k)} I(p_i \in M_k)$$

$$p(M_k) = \frac{1}{3}$$

independently for all doses, and given the 0-1 loss function $\ell(i, M_j)$ in (3) for three decisions, where $i, j \in \{E, S, D\}$, decision rule $\mathcal{D}_{\text{mTPI}}$ in (2) is optimal in the sense that it minimizes the posterior expected loss.

$$\ell(a = i, m_d = M_j) = \begin{cases} 1, & \text{if } i \neq j; \\ 0, & \text{if } i = j, \end{cases} \quad \text{for } i, j \in \{E, S, D\}. \quad (3)$$

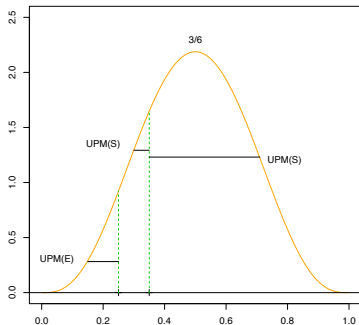
UPM is the marginal posterior probability of each model (interval)

Turns out

$$UPM(k, i) = Pr(M_k \text{ is true} \mid \{x_i, n_i\})$$

This is the marginal posterior probability of model k .

The size of the model is the length of the interval. Ockham's razor picks model $M_S : p_i \in (0.25, 0.35)$; mTPI selects decision "S" when $x_i = 3$ out of $n_i = 6$ patients experience the DLT.

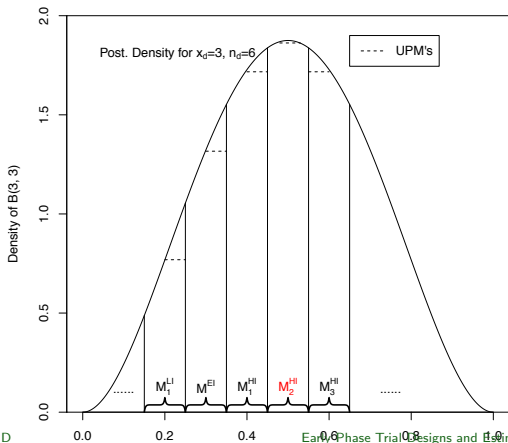


The $mTPI-2$ design (Guo et al., 2017): Blunt the Ockham's Razor

Divid $(0, 1)$ into subintervals with equal length, same as that of $(p_T - \epsilon_1, p_T + \epsilon_2)$.

Pick the decision $\{D, S, E\}$ corresponding to the interval with the largest UPM.

Still the Bayes (optimal) rule



The mTPI-2 Design Compares Favorable to Other Designs

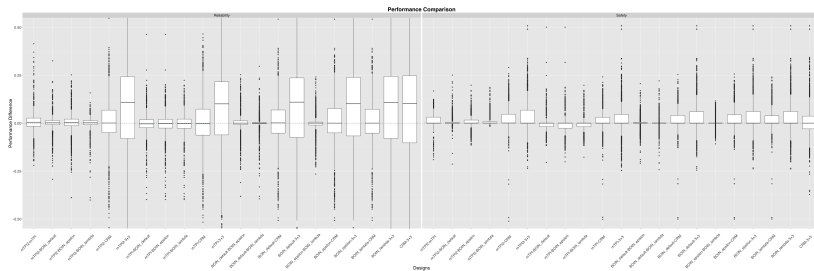
Number of Patients

Number of DLTs	1		2		3		4		5		6		7		8		9		10	
	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2	mTPI	mTPI2
	0	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E
1	D	D	S	D	S	S	S	S	S	E	E	E	E	E	E	E	E	E	E	E
2			DU	DU	D	D	S	D	S	D	S	S	S	S	S	S	S	E	S	E
3					DU	DU	DU	DU	D	D	S	D	S	D	S	D	S	S	S	S
4							DU	DU	DU	DU	DU	DU	D	D	D	D	S	D	S	D
5									DU	DU	DU	DU	DU	DU	DU	DU	DU	DU	D	D
6											DU	DU	DU	DU	DU	DU	DU	DU	DU	DU
7													DU	DU	DU	DU	DU	DU	DU	DU
8															DU	DU	DU	DU	DU	DU
9																	DU	DU	DU	DU
10																			DU	DU

E: Escalate to the next higher dose; **S**: Stay at the same dose; **D**: De-escalate to the previous lower dose; **DU**: De-escalate to the previous lower dose and the current dose will never be used again in the trial;

The mTPI-2 Design Compares Favorable to Other Designs

We compared mTPI, mTPI-2, 3+3, CRM, BOIN in terms of reliability (power of finding MTD) and safety.



Ji and Yang (2017; <https://arxiv.org/abs/1706.03277>)

A New Web-based Integrative Dose-Finding Tool

<http://udesign.laiyaconsulting.com>

A New Web-based Integrative Dose-Finding Tool

<http://udesign.laiyaconsulting.com>

Web Based: No need to download any software; works on MAC, PC, iPad, and smart phones – just need an internet browser (e.g., Chrome, FireFox)

Integrative: Offers up to six designs, 3+3, CRM, mTPI-2, BLRM, etc.

Many new features: CRM decision table, etc.

User-friendly: Demo...

Go beyond toxicity probability intervals

- Interval designs are transparent, simple, and easy to implement. And they are model based.
- Existing drug development more often require **incorporation of efficacy outcome** in dose finding
- Interval designs can be effective for some cases, such as the use of **toxicity and efficacy probability intervals (TEPI)** in Li et al., (2016) for CAR-T therapies
- The posterior probabilities of the intervals can be used to assess the uncertainties of the dose-finding decisions, which can lead to randomized dose-finding designs (ongoing work).

Thank You!