

An overview of modern dose-finding designs for Phase I clinical trials: Part I

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9:30 – 10:10 Part I:
Introduction to Dose-Escalation Trials
The truth about “3+3” design
Model-based Dose-Finding for Single-Agent Trials

10:40 – 11:20 Part II:
Dose-Finding in Health Volunteers
Dose Finding Design for Combination Phase I Trials

Medical & Pharmaceutical Stats (MPS) Research Unit

MPS develops and evaluates novel methods of study design & data analysis for use in the pharmaceutical and medical research community.

We offer number of services:

- Advice on design and analysis of clinical trials
- Develop novel methods for clinical & pre-clinical studies
- Professional Development Courses
 - Design and Analysis of Bioequivalence Studies
 - Pharmacological Modelling
 - Survival and Event History Analysis
 - Adaptive Methods in Clinical Research
 - (!) Designing Early Phase Studies
 - (!) Dose-Finding Designs for Combination Trials

Success rates

According to a recent review (Wong, Siah & Lo, Biostatistics, 2019), between 2000 and 2015

- **41.0%** of confirmatory clinical trials overall and
- **64.5%** of confirmatory clinical trials in oncology

have been unsuccessful.

Reasons for failed confirmatory trials

One of the reasons for failed confirmatory trials are thought to be:

- taking forward treatments that should have been abandoned during early efficacy studies due to insufficient precision when;
 - determining the maximum tolerated dose (MTD);
 - assessing safety;
 - determining the optimal dose.

Consequences

- Avoid going straight into large and expensive Phase III;
- Take more care during Phase I and Phase II trials.

Introduction to Phase I trials

- First experimentation of a new drug in humans
- The emphasis is on **safety**
- Trials are small, typically 20-50 patients
- Patients are added sequentially after side-effects from previous patients have been assessed

Introduction to Phase I trials

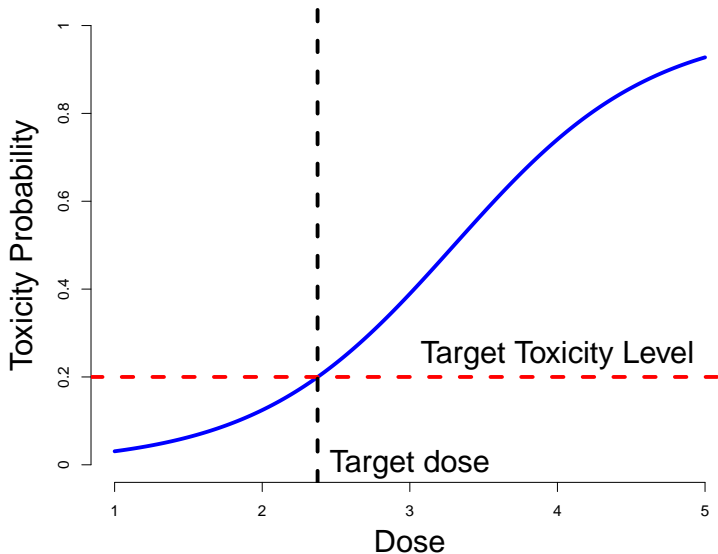
- Subjects
 - **Healthy volunteers** for relatively non-toxic agents
 - **Patients** when drugs are toxic (e.g. in cancer)
- Aim: Find the highest dose with acceptable level of toxicity
 - This is known as the **maximum tolerated dose** (MTD)
 - Based on the assumption that **both benefit** (efficacy) **and risks** (toxicity) of treatment **increase with the dose**
- Setting (of Part I):
 - Binary toxicity outcome (e.g. a *dose-limiting toxicity* (DLT))
 - A *target toxicity level* (TTL) (the desired toxicity at the MTD)

Seeking a quantile

MTD – maximal dose acceptably tolerated by a particular patient population
→ vague

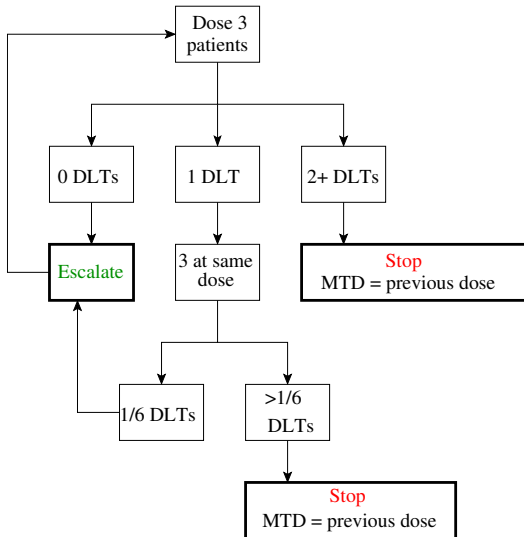
TD 100θ – dose at which the probability of toxicity is θ
(for $0 < \theta < 1$), e.g. TD20
→ more specific

TD20



3+3 design with escalation only

Storer (1989)



- Dose Limiting Toxicity (DLT)
- Simple rule based approach
- No need for a statistician
- Actual dose not used
- The data to declare an MTD are either 0/3 or 1/6

The truth about the 3+3 design

- Example with 4 doses. True toxicities: (0.04, 0.29, 0.36, 0.74)
 - The percentage of patients experimented on each dose are (35%, 43%, 17%, 5%) —**averaged over all possible trials**
 - The recommended MTD probabilities are (48%, 31%, 19%, 0%), 2% no recommended doses
- The 3+3 design
 - tends to underestimate the MTD
 - is inflexible and **memoryless**
 - According to a recent study by Conaway & Petroni (2019), the 3+3 design leads to a **up to 10% noticeably lower success rates** in a Phase III trial compared to model-based alternatives.

A web application for A+B designs:

<https://graham-wheeler.shinyapps.io/AplusB/>

Single-Agent Dose-Finding Study

k increasing doses : $d_1 < d_2 < \dots < d_k$

Response: $x = \begin{cases} 1 & \text{if a patient experienced a DLT} \\ 0 & \text{otherwise} \end{cases}$

Structure: treat successive cohorts of c subjects

Objective: find the “highest safe dose”

Based on the monotonicity assumption: “**the more the better**”:

Both toxicity and efficacy increase with the dose.

Designs for in-patients Phase I trials

Three classes:

- Rule-based designs (e.g. “3+3” design)
- Model-based designs (e.g. CRM, EWOC, etc.)
- Model-assisted (shape-free) designs (mTPI, BOIN, etc.)

Review of advantages Jaki *et al.* (2013)

General (Bayesian) model-based design

Before the trial:

- 1 Choose doses d_1, \dots, d_k ;
- 2 Choose a form of dose-response relationship $p(d_i, \alpha)$ where α are model parameters;
- 3 Impose a prior distribution for α ;
- 4 Choose a criterion to allocate patients;
- 5 Choose stopping rules (e.g. estimated accurately enough).

During the trial:

- 1 Sequentially update estimates of α ;
- 2 Select the dose for the next cohort using the criterion;
- 3 Stop if at least one of the stopping rules is met.

General (Bayesian) model-based design

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Continual Reassessment Method (CRM)

by O'Quigley et al (1990)

Response: Binary

Model: $p(\pi_i, \alpha) = \pi_i^{\exp(\alpha)}$, π_i are standardised doses

(skeleton) calculated from prior estimates of p_i

Prior on α : Normal $\alpha \sim \mathcal{N}(\mu, \sigma^2)$

Allocation Rule: $\min |p_i(\pi_i, \hat{\alpha}) - \theta|$ where $\hat{\alpha} = \mathbb{E}(\alpha)$

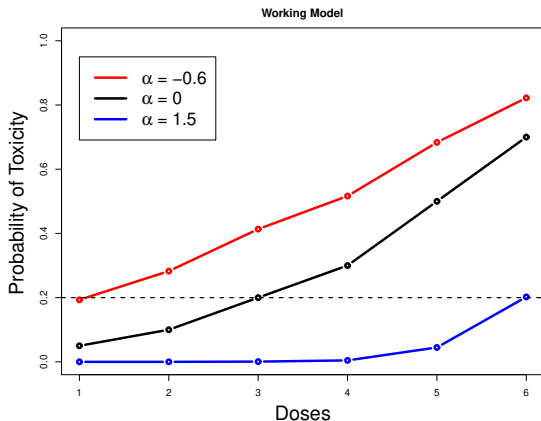
This form of the CRM is extensively studied in the literature:

- Prior for α : $\mathcal{N}(0, 1.34)$ → all doses has the same prior probability to be the MTD
- Operational skeleton: given “equivalent interval” → an optimal spacing between skeleton

Representation of the model

Starting values for π_i

π_1	π_2	π_3	π_4	π_5	π_6
0.05	0.10	0.20	0.30	0.50	0.70



Bayesian updating

- 1 Specify the prior distribution of α
- 2 Assign the first cohort to the lowest dose
- 3 Given the observations (data) and the prior distribution, update the (posterior) distribution of α

$$\text{Posterior} \propto \text{Prior} \times \text{Data}$$

- 4 Given the posterior find the “best” guess of α : $\hat{\alpha}$
- 5 Find estimates of toxicity probabilities as $\hat{p}_i = \pi_i^{\exp(\hat{\alpha})}$
- 6 Allocate the next cohort to the dose having the estimated toxicity closest to the target level θ .
- 7 Repeat steps 4-6 using the obtained Posterior as Prior.

Alternative dose-toxicity model

Model:

- Two-parameter logistic regression model

$$p(d_{(j)}) = \frac{\exp\{\alpha_1 + \alpha_2 \log(d_{(j)})\}}{1 + \exp\{\alpha_1 + \alpha_2 \log(d_{(j)})\}}$$

- where $d_{(1)} < \dots < d_{(k)}$ are doses (!)

Requires a **prior** distribution on (α_1, α_2)

Similar to the one-parameter CRM, there is a recommendation for this prior that leads to good operating characteristics

$$(\log \alpha_1, \log \alpha_2) \sim \mathcal{N} \left(\begin{bmatrix} 2.15 \\ 0.52 \end{bmatrix}, \begin{bmatrix} 0.84^2 & 0.134 \\ 0.134 & 0.80^2 \end{bmatrix} \right)$$

Allocation Criteria

- Escalation with Overdose Control (EWOC):

$$\mathbb{E} (\nu(\theta - p_i)^+ + (1 - \nu)(p_i - \theta)^+)$$

e.g. $\nu = 0.25$

- NCRM by Neuenschwander et al (2008):
 - Maximising the probability of being in the target interval
 - while safeguarding the patients (controlling the probability that dose is too toxic)

Comments on the implementation

Planning and conducting the trial using model-based designs

- Model-based designs would required more effort to be implemented. However, there is a variety of software implementing these designs.
- There are several ready-to-use interactive Web Applications for the designs covered above that require **no programming/statistical skills**

They can be used for **simulation and implementation** of different model-based designs

- 1-parameter CRM `uvatrapps.shinyapps.io/crmb/`
- 2-parameter + prior elicitation `lancs.shinyapps.io/Design/`



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