



# Taking your agents to the clinic: phase I study design

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# Dose finding

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- Traditional goal: Find the highest dose with acceptable toxicity
- New goals:
  - find dose with sufficient effect on biomarker
  - find dose with acceptable toxicity and high efficacy
  - Find dose with acceptable toxicity in the presence of another agent that may also be escalated.

# Acceptable toxicity

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- What is acceptable rate of toxicity?
  - 20%?
  - 30%?
  - 50%?
- What is toxicity????
  - Standard in cancer: Grade 4 hematologic or grade 3/4 non-hematologic toxicity
  - Always?
  - Does it depend on reversibility of toxicity?
  - Does it depend on intensity of treatment?
    - Tamoxifen?
    - Chemotherapy?

# Phase I study design

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- “Standard” Phase I trials (in oncology) use what is often called the ‘3+3’ design (aka ‘modified Fibonacci’):

Treat 3 patients at dose K

1. If 0 patients experience dose-limiting toxicity (DLT), escalate to dose K+1
2. If 2 or more patients experience DLT, de-escalate to level K-1
3. If 1 patient experiences DLT, treat 3 more patients at dose level K
  - A. If 1 of 6 experiences DLT, escalate to dose level K+1
  - B. If 2 or more of 6 experiences DLT, de-escalate to level K-1

- Maximum tolerated dose (MTD) is considered highest dose at which 1 or 0 out of six patients experiences DLT.
- Doses need to be pre-specified
- Confidence in MTD is usually poor.

# Should we use the “3+3”?

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- **It is terribly imprecise and inaccurate in its estimate of the MTD**
- **Why?**
  - **MTD is not based on all of the data**
  - **Algorithm-based method**
  - **Ignores rate of toxicity!!!**
- Likely outcomes:
  - Choose a dose that is too high
    - Find in phase II that agent is too toxic.
    - Abandon further investigation or go back to phase I
  - Choose a dose that is too low
    - Find in phase II that agent is ineffective
    - Abandon agent

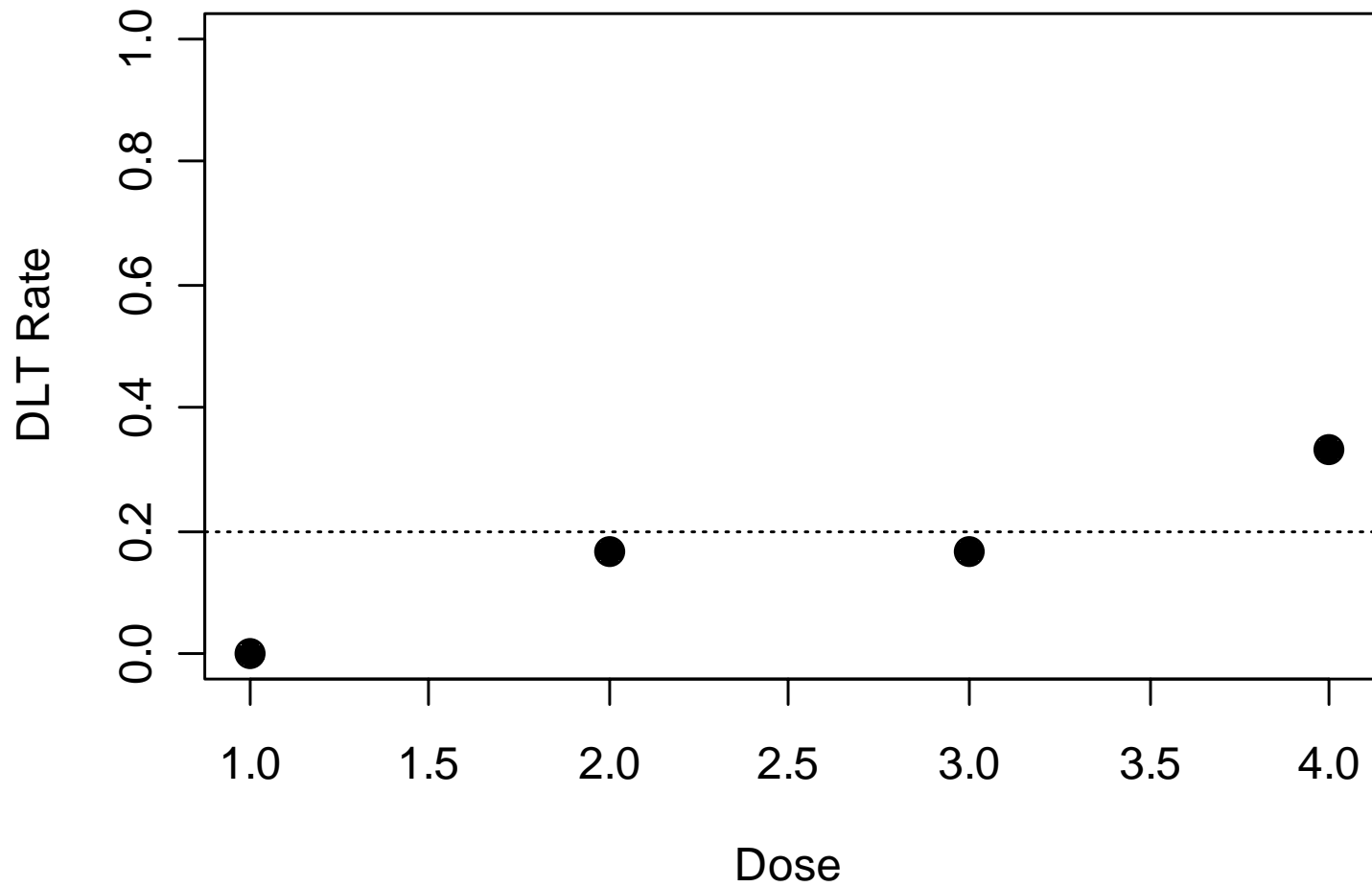
# Two examples:

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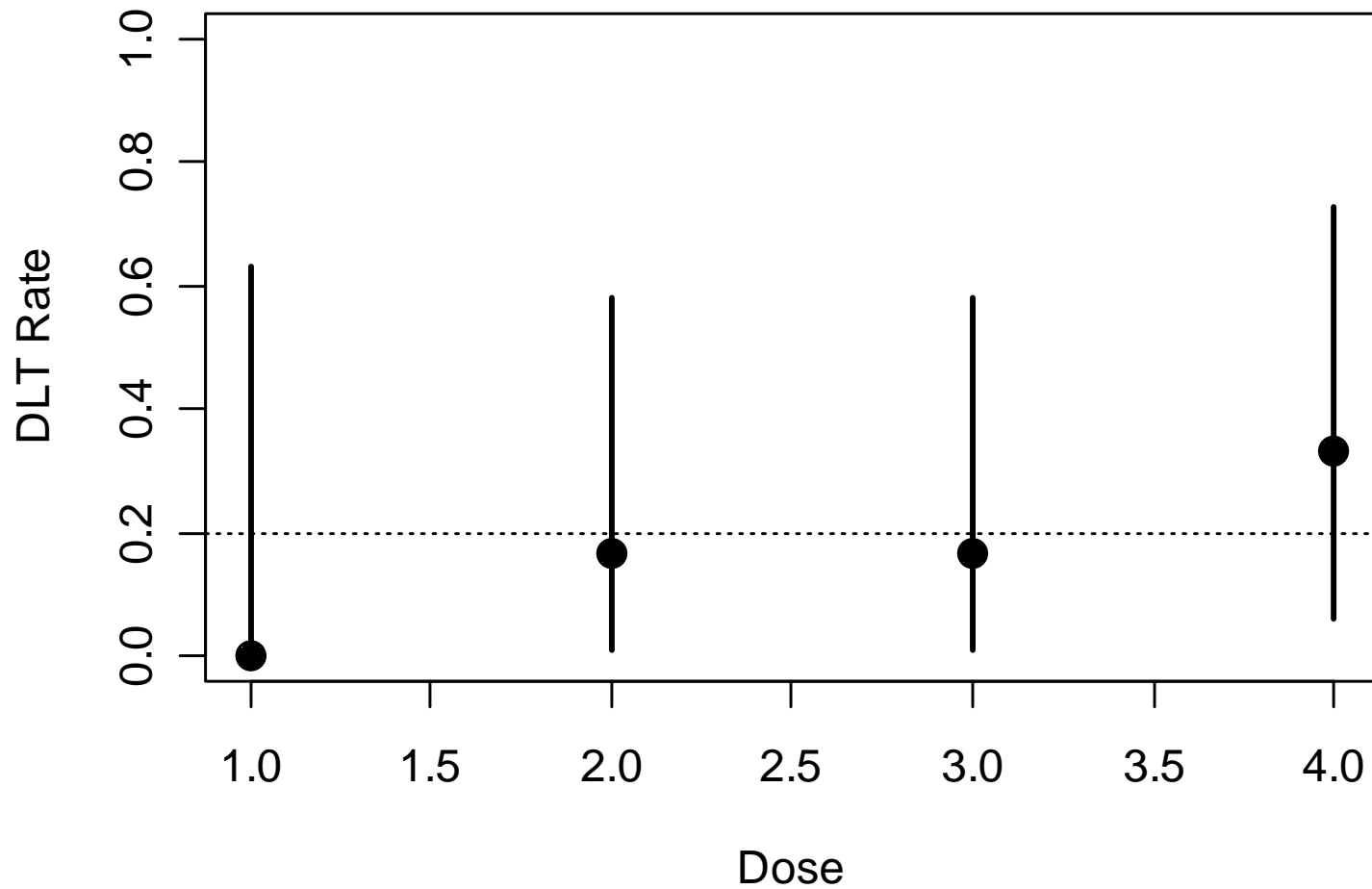
Example 1: total N=21

	Cohort 1	Cohort 2	Cohort 3	Cohort 4	Cohort 5	Cohort 6	Cohort 7
Dose	1	2	2	3	3	4	4
DLTs	0/3	1/3	0/3	1/3	0/3	1/3	1/3

# Observed Data



# Observed Data: with 90% CIs





# Example 2:

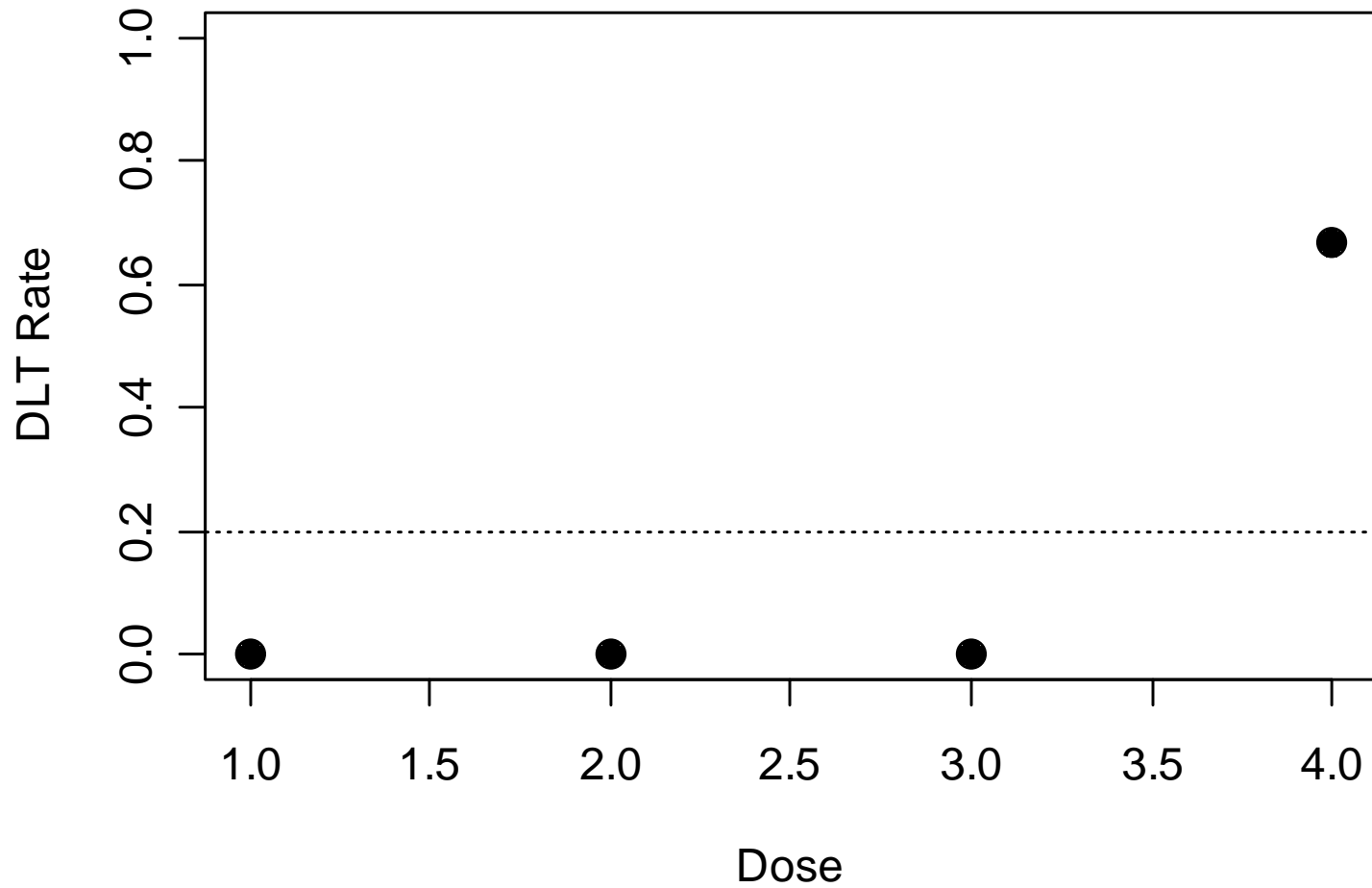
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Example 2: total N=12

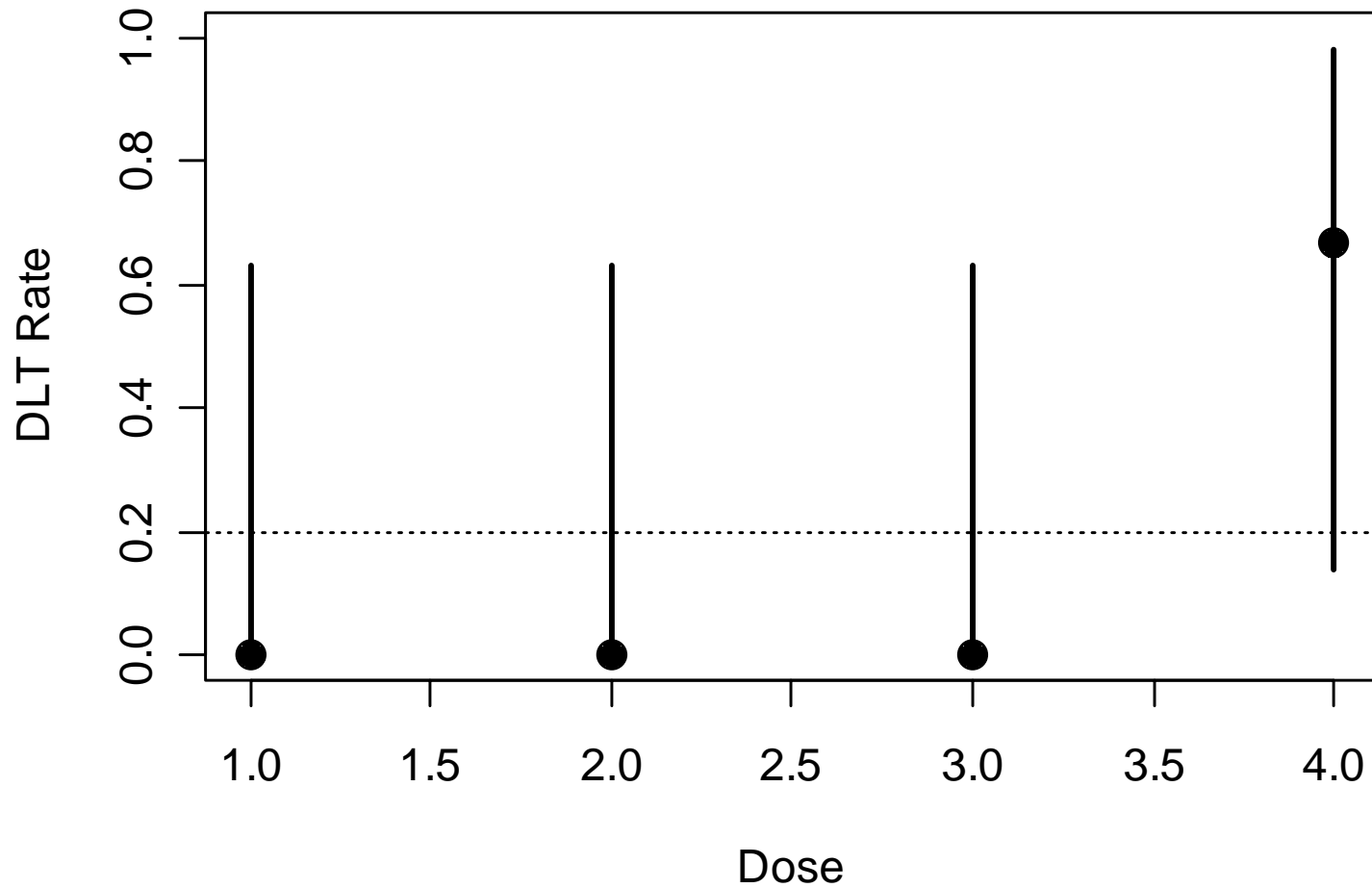
	Cohort 1	Cohort 2	Cohort 3	Cohort 4
Dose	1	2	3	4
DLTs	0/3	0/3	0/3	2/3

# Observed Data

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# Observed Data: with 90% CIs



# Why is the 3+3 so popular?

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- People know how to implement it
- “we just want a quick phase I”
- It has historic presence
- FDA (et al.) accept and promote it
- There is a level of comfort from the approach
- The “better” approaches are too statistical(!)

# USE A SMARTER DESIGN!

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- **Phase I is the most critical phase of drug development!**
- **What makes a good design?**
  - Accurate selection of MTD
    - dose close to true MTD
    - dose has DLT rate close to the one specified
  - Relatively few patients in trial are exposed to toxic doses
- **Why not impose a statistical model?**
- **What do we “know” that would help?**
  - **Monotonicity**
  - **Desired level of DLT**

# Continual Reassessment Method (CRM)

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- Allows statistical modeling of optimal dose: dose-response relationship is assumed to behave in a certain way
- Can be based on “safety” or “efficacy” outcome (or both).
- Design searches for best dose **given a desired toxicity or efficacy level** and does so in an efficient way.
- **This design REALLY requires a statistician throughout the trial.**
- **ADAPTIVE**

# CRM history in brief

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- Originally devised by O'Quigley, Pepe and Fisher (1990) **where dose for next patient was determined based on responses of patients previously treated in the trial**
- Due to safety concerns, several authors developed variants
  - Modified CRM (Goodman et al. 1995)
  - Extended CRM [2 stage] (Moller, 1995)
  - Restricted CRM (Moller, 1995)
  - and others....

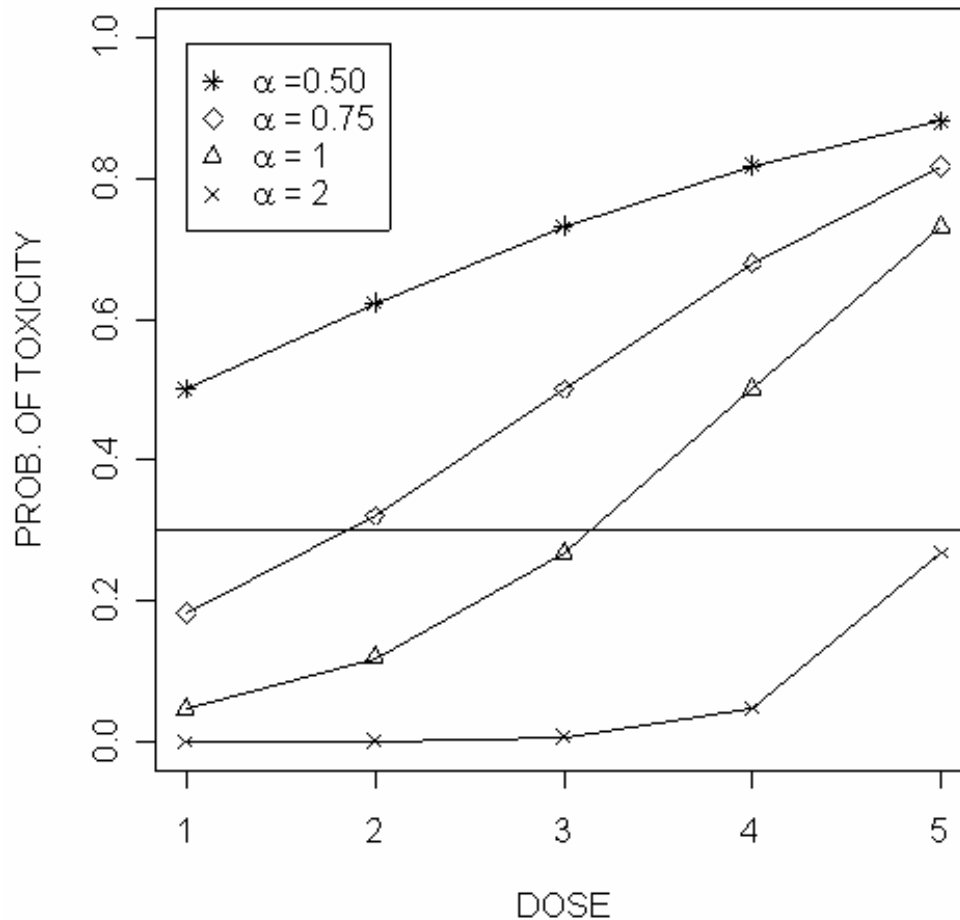
# Some reasons why to use CRM

Table 4 Characteristics of five Phase 1 studies. Designs 1–3 are CRM designs with different sample sizes and cohort sizes. Designs 4–6 are “3 + 3” designs with different prespecified dose levels. All six designs have the same true model of dose–toxicity. Dose levels for the “3 + 3” designs are shown in Figure 5. Results are based on 1000 simulated trials for each design

	Design 1 (CRM) (Example 2)	Design 2 (CRM)	Design 3 (CRM)	Design 4 ("3 + 3")	Design 5 ("3 + 3")	Design 6 ("3 + 3")
1 Total sample size: fixed for CRMs, median for “3 + 3”	30	50	60	27	27	39
2 Patients per cohort	3	5	3	3	3	3
3 Number of cohorts fixed for CRMs, median for “3 + 3”	10	10	20	9	9	13
4 % of trials with recommended dose within 250mg of true dose (1656 mg)	57%	72%	71%	41%	35%	32%
5 % of trials with recommended dose within 400mg of true dose (1656 mg)	80%	91%	89%	41%	54%	58%
6 % of trials with recommended dose dose-limiting toxicity rate of >40%	9.5%	5.8%	5.9%	7.1%	21%	12%
7 % of trials with recommended dose dose-limiting toxicity rate of >50%	0.9%	0.2%	0.6%	7.1%	2.0%	0.6%
8 % of trials with recommended dose dose-limiting toxicity rate of <20%	13%	5.7%	6.2%	52%	44%	38%
9 % of trials with recommended dose dose-limiting toxicity rate of <10%	0.0%	0.1%	0.0%	11%	16%	6.9%
10 Average % of patients treated at doses with 40% or greater dose-limiting toxicity rate	7.6%	7.8%	5.7%	17%	23%	7.5%
11 Average % of patients treated at doses with 20% or less dose-limiting toxicity rate	32%	19%	24%	62%	53%	64%
12 Average % of patients with dose-limiting toxicities	26%	28%	26%	21%	22%	19%



# Basic Idea of CRM



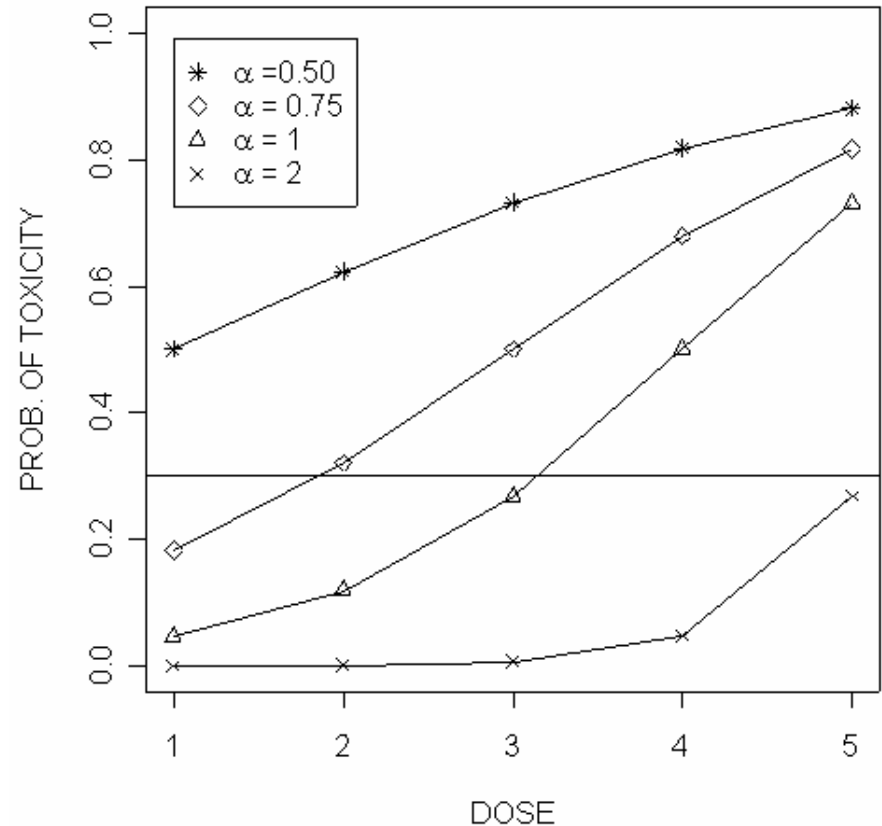
$$p(\text{toxicity} | \text{dose} = d_i) = \frac{\exp(3 + \alpha d_i)}{1 + \exp(3 + \alpha d_i)}$$

# Modified CRM

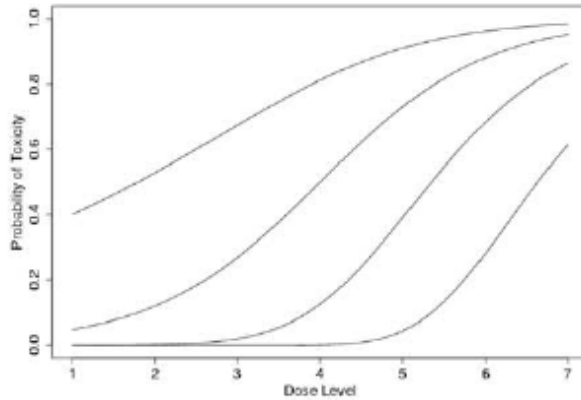
(Goodman, Zahurak, and Piantadosi, *Statistics in Medicine*, 1995)

## Carry-overs from standard CRM

- Mathematical dose-toxicity model must be assumed
- To do this, need to think about the dose-response curve and get preliminary model.
- **We CHOOSE the level of toxicity that we desire for the MTD (e.g.,  $p = 0.30$ )**
- At end of trial, we can estimate dose response curve.

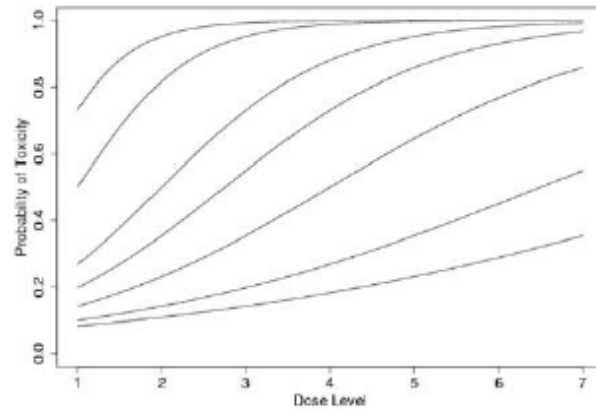


# Some other mathematical models we could choose

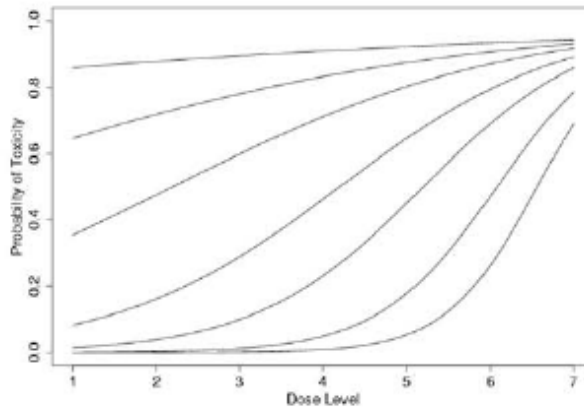


A. Hyperbolic

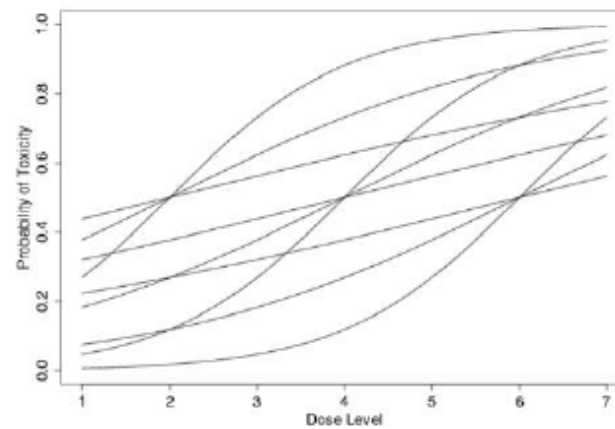
tangent: 
$$p(dose) = \left( \frac{\tanh(dose) + 1}{2} \right)^b$$



B. One-parameter logistic: 
$$p(dose) = \frac{e^{-4+b \cdot dose}}{1 + e^{-4+b \cdot dose}}$$



C. One-parameter logistic: 
$$p(dose) = \frac{e^{3+b \cdot dose}}{1 + e^{3+b \cdot dose}}$$



D. Two-parameter logistic: 
$$P = \frac{e^{a+b \cdot dose}}{1 + e^{a+b \cdot dose}}$$

Modified CRM by  
Goodman, Zahurak, and Piantadosi  
(Statistics in Medicine, 1995)

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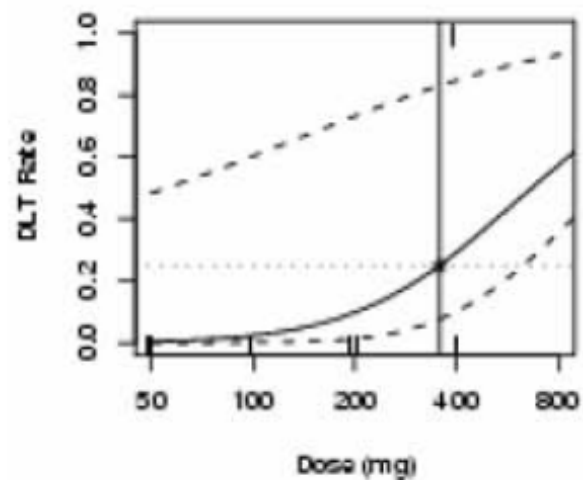
- Modifications by Goodman et al.
  - Use ‘standard’ dose escalation model until first toxicity is observed:
    - Choose cohort sizes of 1, 2, or 3
    - Use standard ‘3+3’ design (or, in this case, ‘2+2’)
  - **Upon first toxicity, fit the dose-response model using observed data**
    - Estimate  $\alpha$
    - Find dose that is closest to desired toxicity rate.
  - Does not allow escalation to increase by more than one dose level.
  - De-escalation can occur by more than one dose level.

# Simulated Example

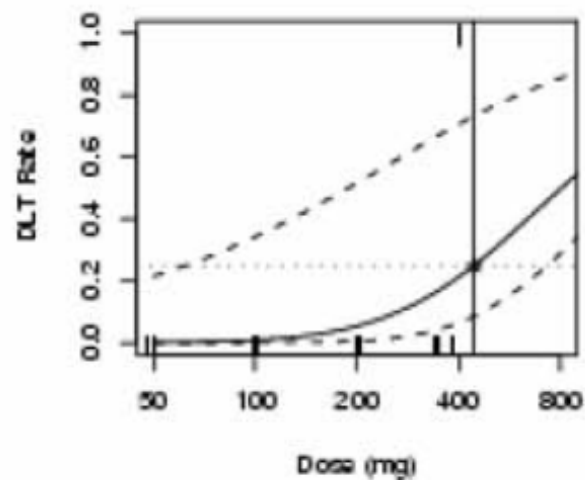
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- Shows how the CRM works in practice
- Assume:
  - Cohorts of size 2
  - Escalate at fixed doses until DLT occurs
  - Then, fit model and use model-based escalation
  - Increments of 50mg are allowed
  - Stop when 10 patients have already been treated at a dose that is the next chosen dose

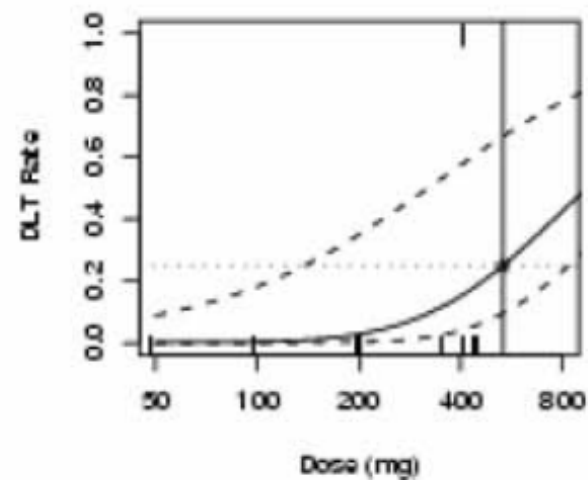
Cohort 4



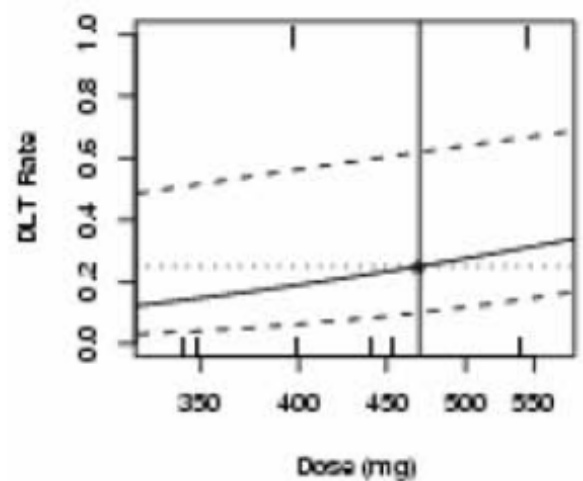
Cohort 5



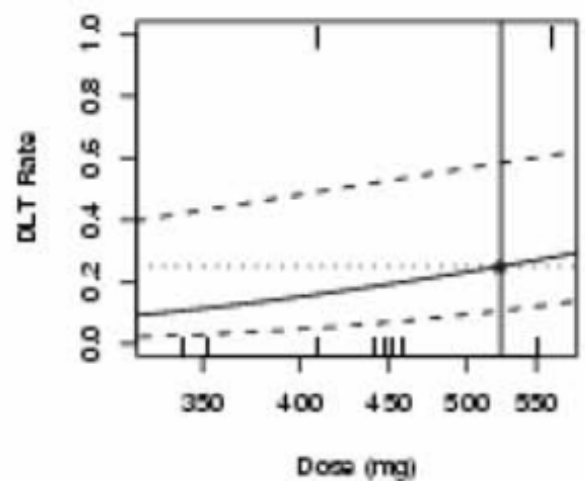
Cohort 6



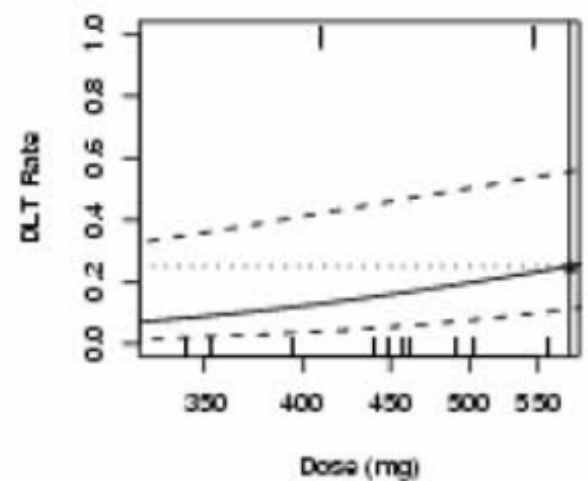
Cohort 7



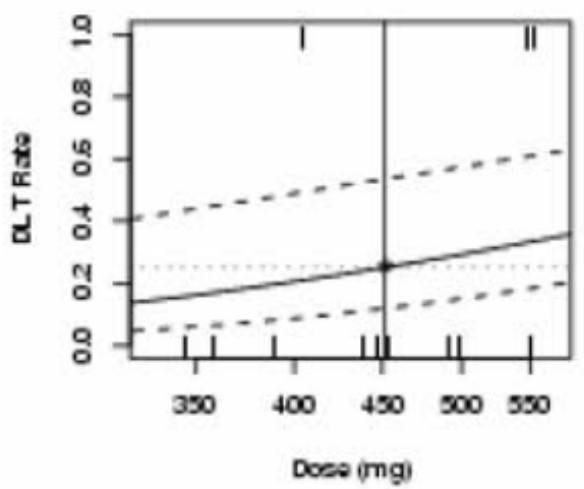
Cohort 8



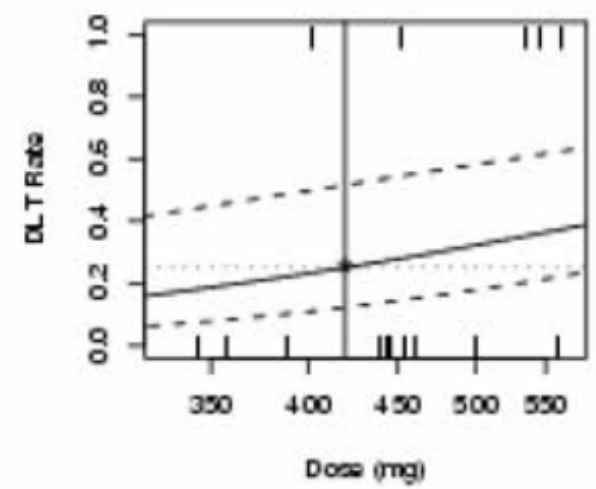
Cohort 9



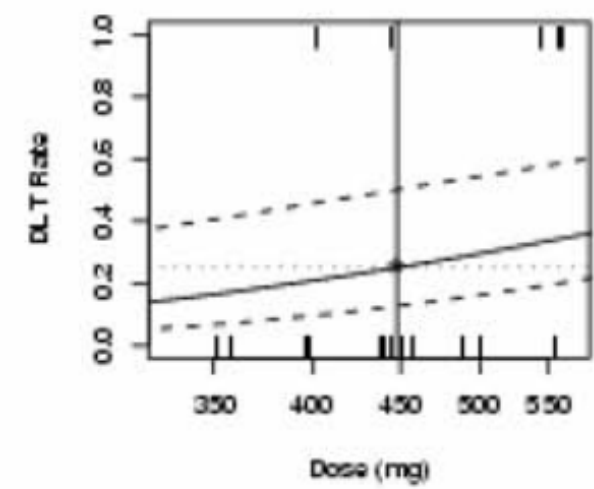
Cohort 10



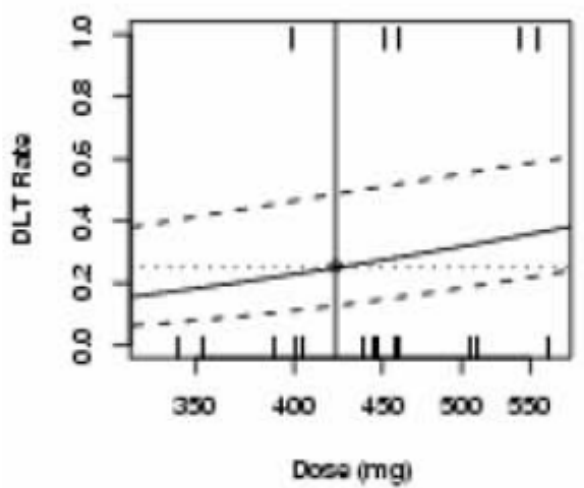
Cohort 11



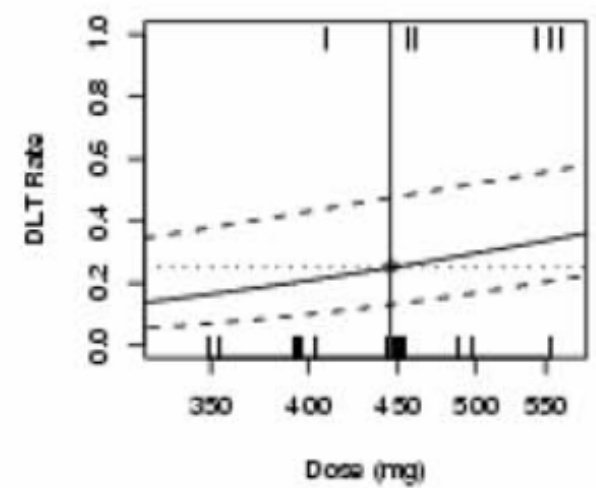
Cohort 12



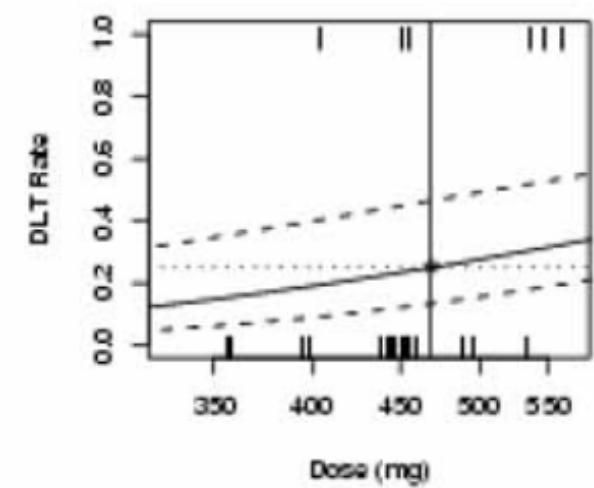
Cohort 13



Cohort 14



Cohort 15



# Result

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- 450mg is determined to be the optimal dose to take to phase II
- 30 patients (?!)
- Confidence interval for true DLT rate at 450mg: 15% - 40%
- Used ALL of the data to make our conclusion



## Real Example Samarium in pediatric osteosarcoma:

**Desired DLT rate is 30%.**

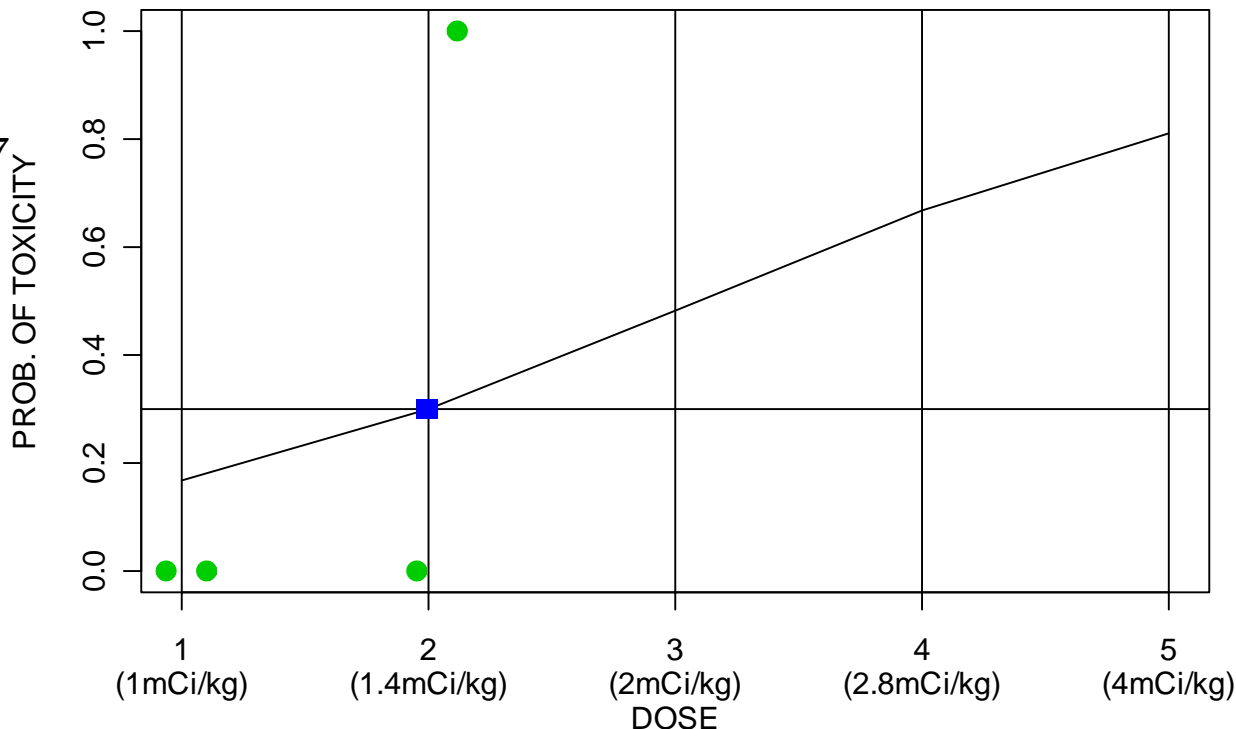
2 patients treated at dose 1 with 0 toxicities

2 patients treated at dose 2 with 1 toxicity

→ Fit CRM using equation below

$$p(\text{toxicity} | \text{dose} = d_i) = \frac{\exp(3 + \alpha d_i)}{1 + \exp(3 + \alpha d_i)}$$

- Estimated  $\alpha = 0.77$
- Estimated dose is 1.4mCi/kg for next cohort.



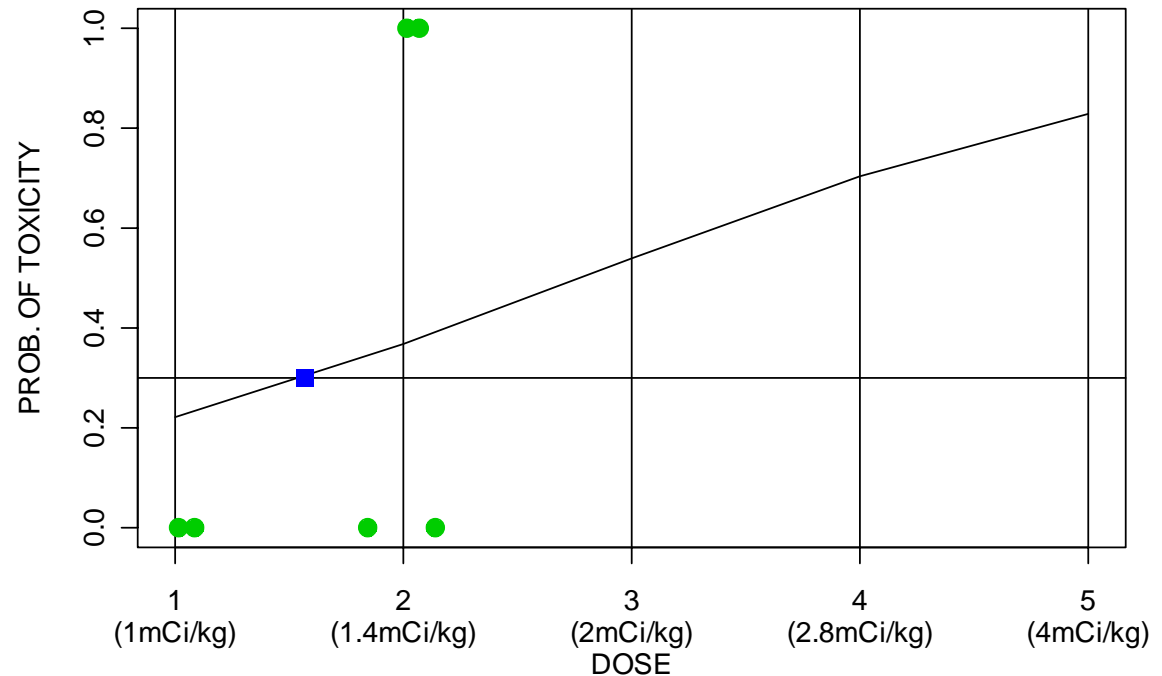
## **Example** Samarium study with cohorts of size 2:

2 patients treated at 1.0 mCi/kg with no toxicities

4 patients treated at 1.4 mCi/kg with 2 toxicities

→ Fit CRM using equation on earlier slide

- Estimated  $\alpha = 0.71$
- Estimated dose for next patient is 1.2 mCi/kg



## **Example** Samarium study with cohorts of size 2:

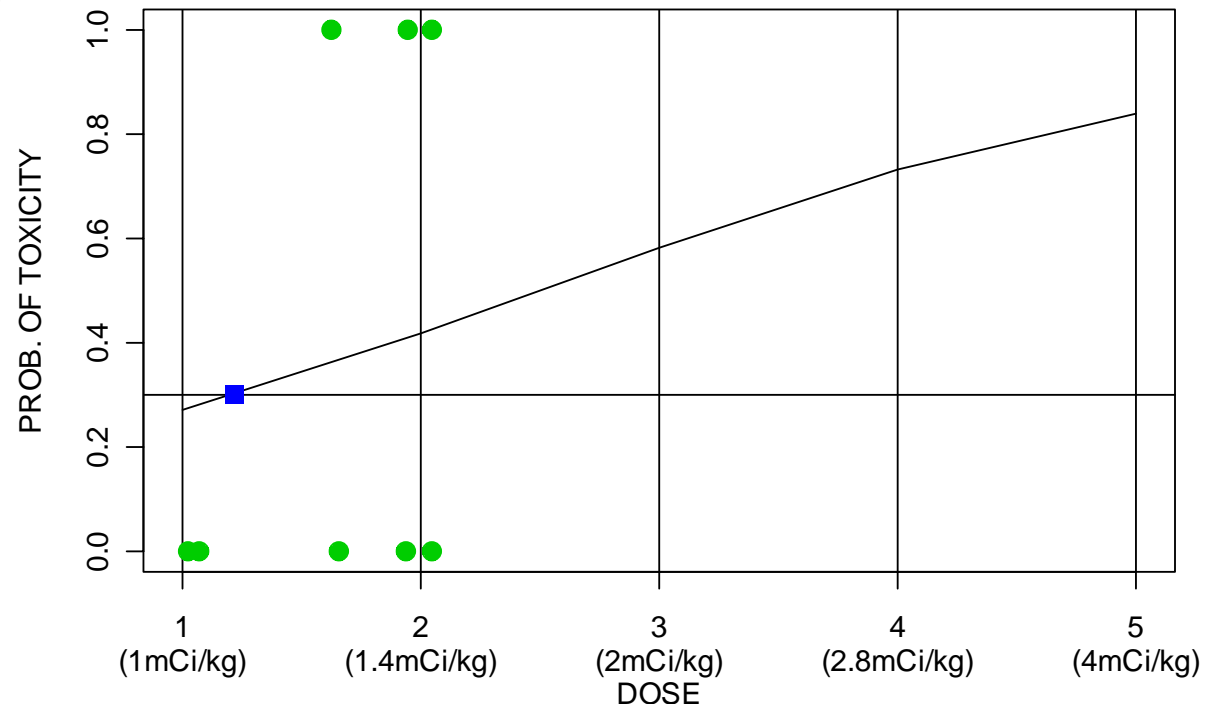
2 patients treated at 1.0 mCi/kg with no toxicities

4 patients treated at 1.4 mCi/kg with 2 toxicities

2 patients treated at 1.2 mCi/kg with 1 toxicity

→ Fit CRM using equation on earlier slide

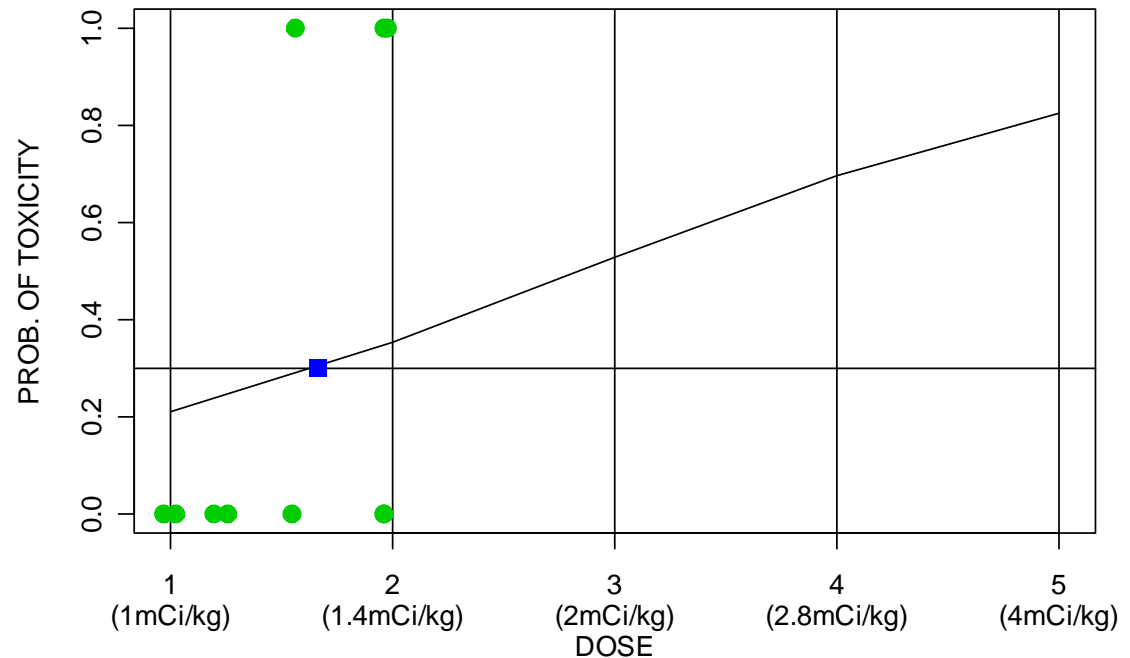
- Estimated  $\alpha = 0.66$
- Estimated dose for next patient is 1.1 mCi/kg



## **Example** Samarium study with cohorts of size 2:

- 2 patients treated at 1.0 mCi/kg with no toxicities
  - 4 patients treated at 1.4 mCi/kg with 2 toxicities
  - 2 patients treated at 1.2 mCi/kg with 1 toxicity
  - 2 patients treated at 1.1 mCi/kg with no toxicities
- Fit CRM using equation on earlier slide

- Estimated  $\alpha = 0.72$
- Estimated dose for next patient is 1.2 mCi/kg



# When does it end?

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- Pre-specified stopping rule
- Can be fixed sample size
- Often when a “large” number have been assigned to one dose.
- This study should enroll at least two more cohorts.

# Dose increments

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- Can be discrete or continuous
- Infusion?
- Tablet?
- Stopping rule should depend on nature (and size) of allowed increment!

# A little more on the statistics:

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- Original design was purely Bayesian
- Requires a prior distribution
  - Prior is critically important because it outweighs the data early in the trial
  - Computationally is somewhat challenging
- Some revised designs use ML
  - Simpler to use
  - Once a DLT is observed, model can be fit
  - Some will “inform” the ML approach using “pseudo-data” (Piantadosi)

# Simple prediction, but backwards(?)

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- Usual prediction:
  - Get some data
  - Fit model
  - Estimate the **outcome** for a new patient with a particular **characteristic**
- CRM prediction
  - Get some data
  - Fit model
  - Find the **characteristic** (dose) associated with a particular **outcome** (DLT rate)



# Finding the next dose: ML approach

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- Use maximum likelihood to estimate the model.
- What likelihood do we use? Binomial.

$$L(p; y) = \prod_{i=1}^N p^{y_i} (1-p)^{(1-y_i)}$$

$$p(\text{toxicity} | \text{dose} = d_i) = \frac{\exp(3 + \alpha d_i)}{1 + \exp(3 + \alpha d_i)}$$

- Algorithmic estimation of  $\alpha$

# Finding next dose

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- Recall model, now with estimated  $\alpha$ :

$$p_i = \frac{\exp(3 + \hat{\alpha}d_i)}{1 + \exp(3 + \hat{\alpha}d_i)}$$

- Rewrite in terms of  $d_i$ :

$$d_i = \frac{\log\left(\frac{p_i}{1-p_i}\right) - 3}{\hat{\alpha}}$$

# Finding next dose

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- Use desired DLT rate as  $p_i$

$$d_i = \frac{\log(\frac{3}{7}) - 3}{\hat{\alpha}} = \frac{-3.85}{\hat{\alpha}}$$

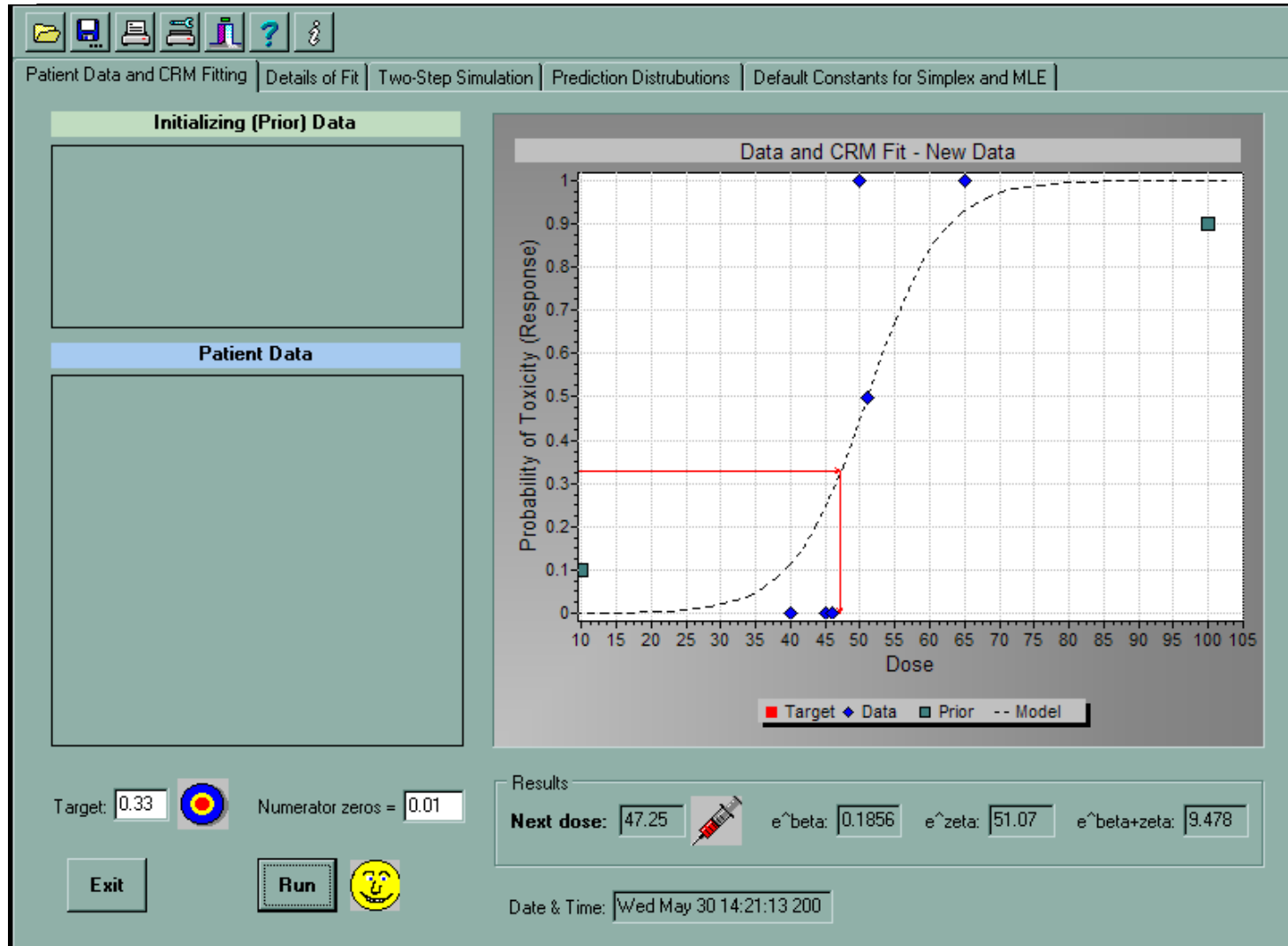
# Negative dose?

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- Doses are often mapped to another scale
- dose coding:
  - 6 = level 1 (1.0)
  - 5 = level 2 (1.4)
  - 4 = level 3 (2.0)
  - 3 = level 4 (2.8)
  - 2 = level 5 (4.0)
- WHY? Makes the statistics work....

# CRM Software:

<http://www.cancerbiostats.onc.jhmi.edu/software.cfm>



# Escalation with Overdose Control

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- EWOC (Babb et al.)
- Similar to CRM
- Bayesian
- Advantage: overdose control
  - “loss function”
  - Constrained so that the predicted proportion of patients who receive an overdose cannot exceed a specified value
  - **Implies that giving an overdose is greater mistake than an underdose**
  - CRM does not make this distinction
  - This control is changed as data accumulates

Figure 3. Proportion of patients given doses for which the probability of a severe toxic reaction is less than or equal to 1/5. Each × and + represents the results from all simulation runs for a particular parameter combination. Each point, (●), is the average of the results obtained for a particular method at the six parameter combinations considered. For each of the UD schemes the results obtained when only 6 dose levels were used are shown to the left of the results obtained when 11 levels were used

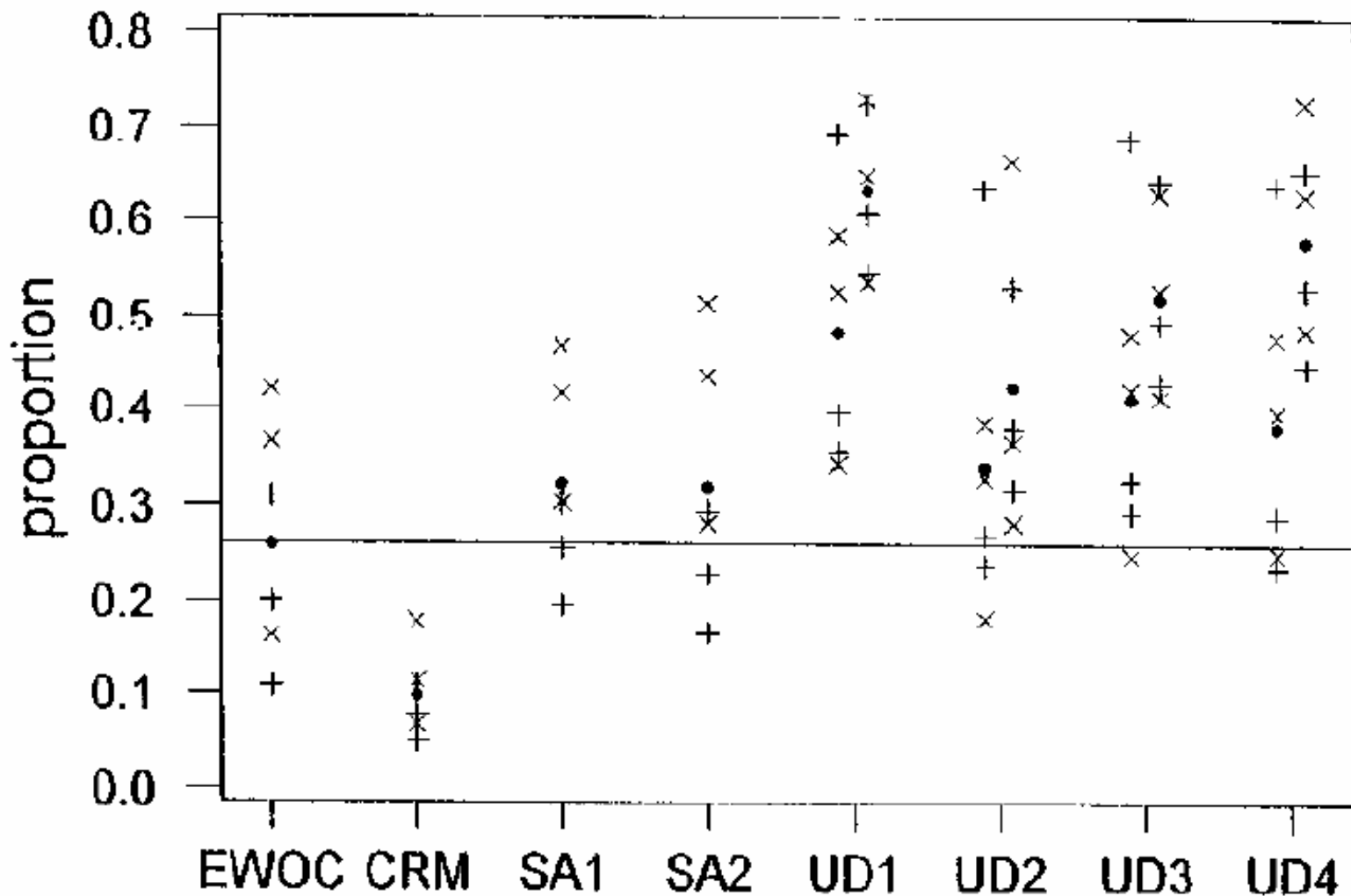


Figure 9. Average bias of  $\hat{\gamma}$ , the estimate of the MTD

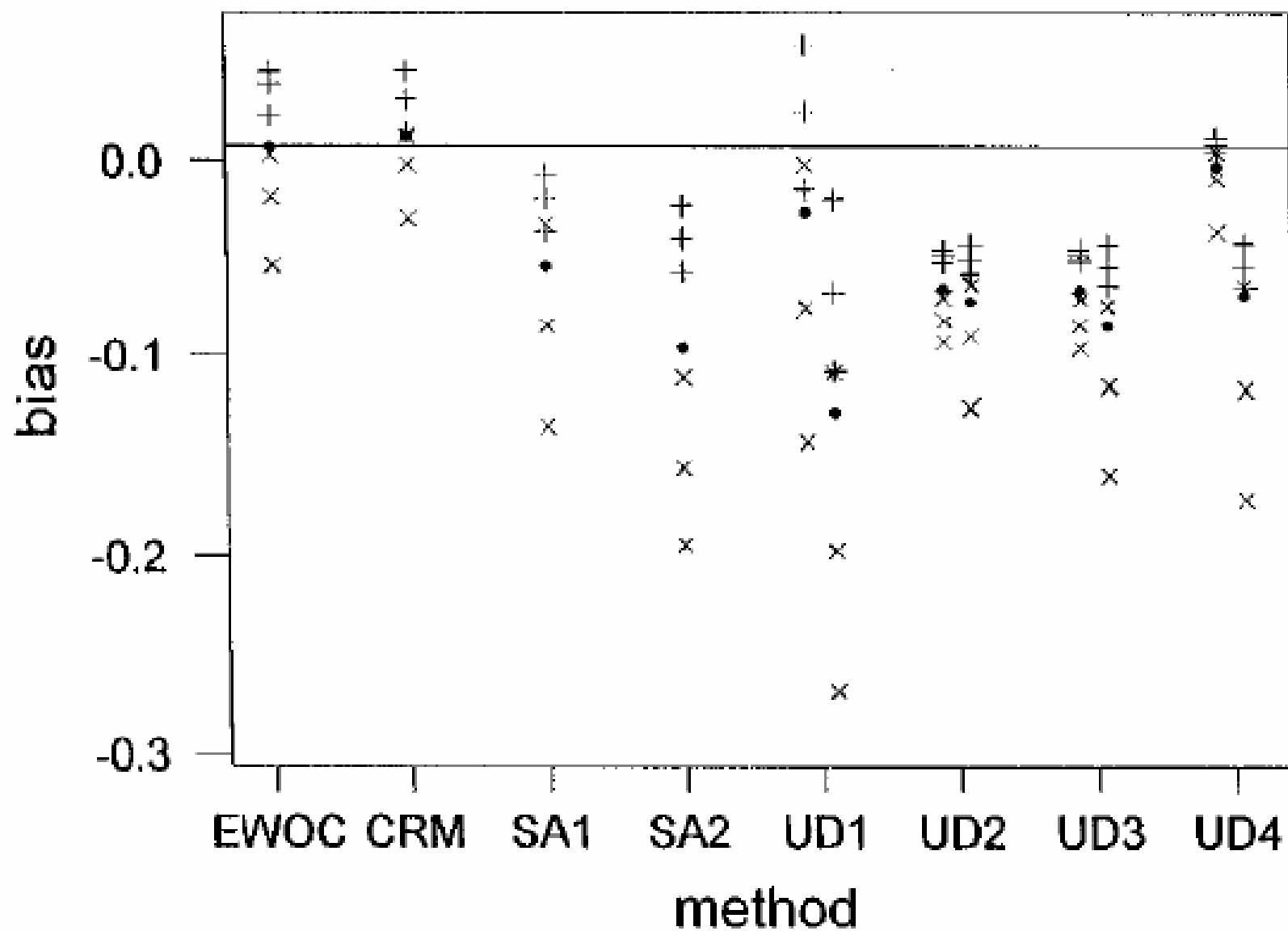
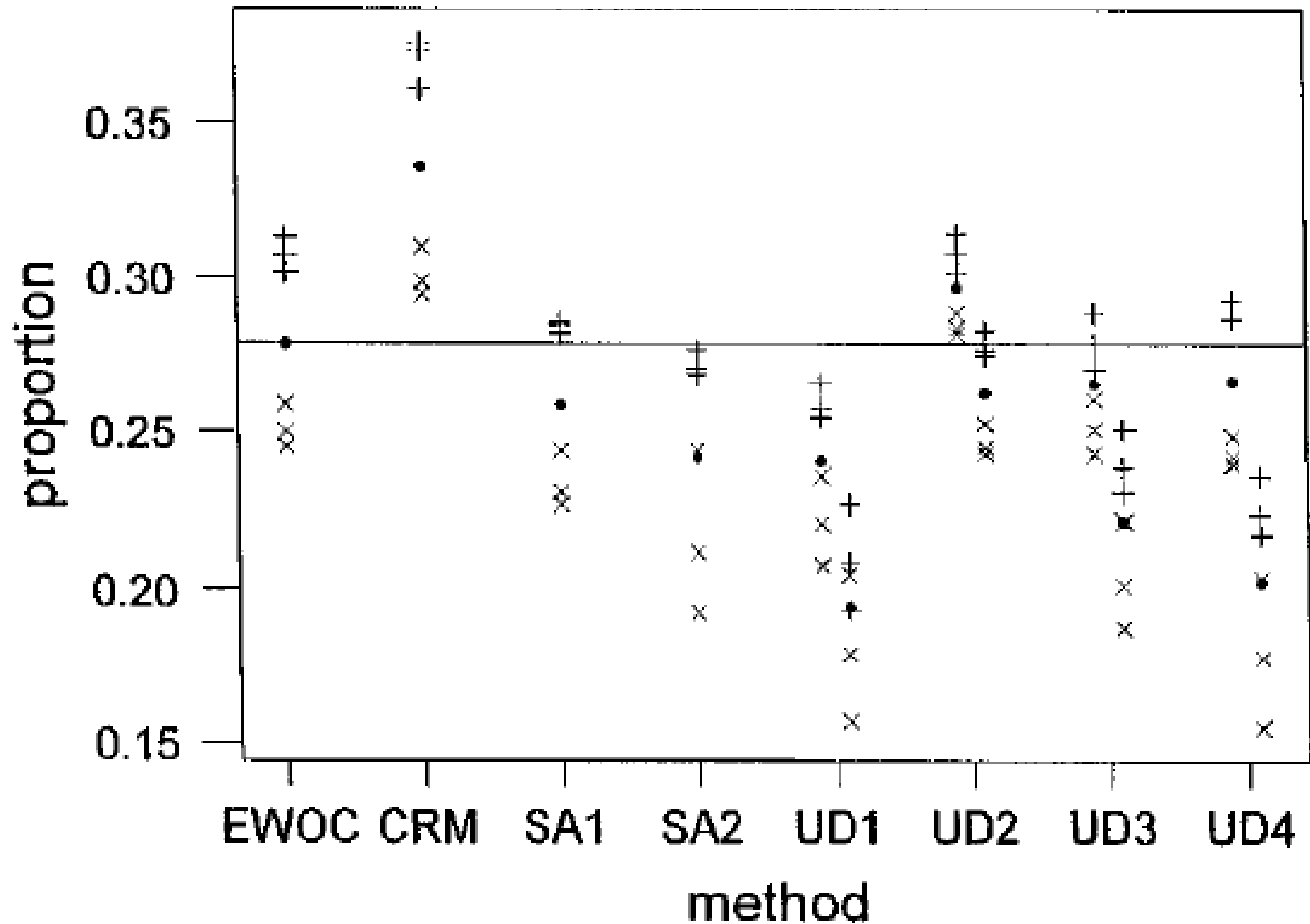


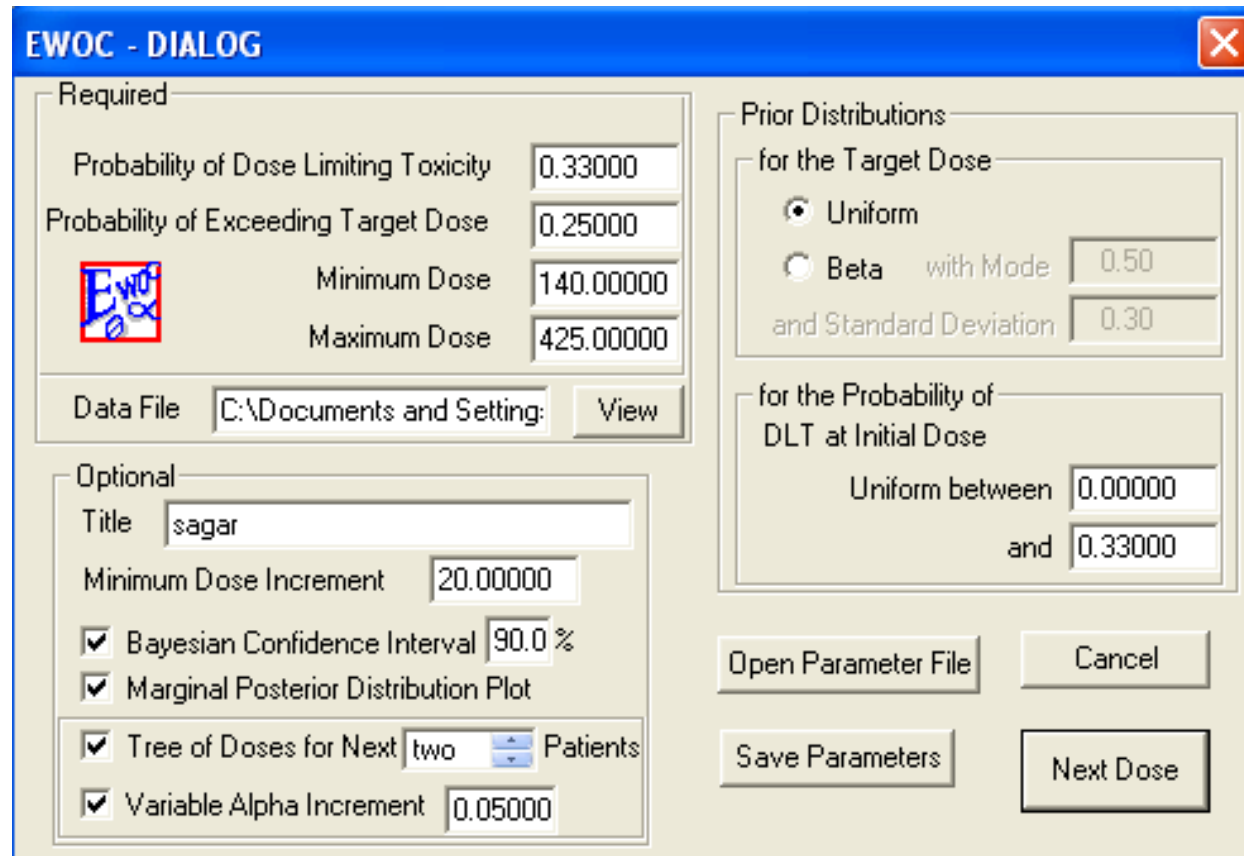


Figure 12. Proportion of patients exhibiting dose-limiting toxicity



# EWOC Software

- <http://www.sph.emory.edu/BRI-WCI/ewoc.html>



The image shows a screenshot of the EWOC - DIALOG software interface. The window title is "EWOC - DIALOG" and it has a standard Windows-style title bar with a close button. The interface is divided into several sections:

- Required:** This section contains four input fields: "Probability of Dose Limiting Toxicity" (0.33000), "Probability of Exceeding Target Dose" (0.25000), "Minimum Dose" (140.00000), and "Maximum Dose" (425.00000). There is also a small logo with the letters "EWOC" and a blue circle.
- Data File:** A text box shows the path "C:\Documents and Setting:" followed by a "View" button.
- Optional:** This section includes a "Title" field with the text "sagar", a "Minimum Dose Increment" field (20.00000), and four checked checkboxes: "Bayesian Confidence Interval" (90.0%), "Marginal Posterior Distribution Plot", "Tree of Doses for Next" (two Patients), and "Variable Alpha Increment" (0.05000).
- Prior Distributions:** This section is divided into two sub-sections. The first, "for the Target Dose", has two radio buttons: "Uniform" (selected) and "Beta" (with Mode 0.50 and Standard Deviation 0.30). The second, "for the Probability of DLT at Initial Dose", has a "Uniform between" field (0.00000) and an "and" field (0.33000).

At the bottom right, there are four buttons: "Open Parameter File", "Cancel", "Save Parameters", and "Next Dose".



# Other Novel Ideas in Phase I

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- Outcome is not always toxicity
- Even in phase I, efficacy can be outcome to guide dose selection
- Two outcomes: safety and efficacy

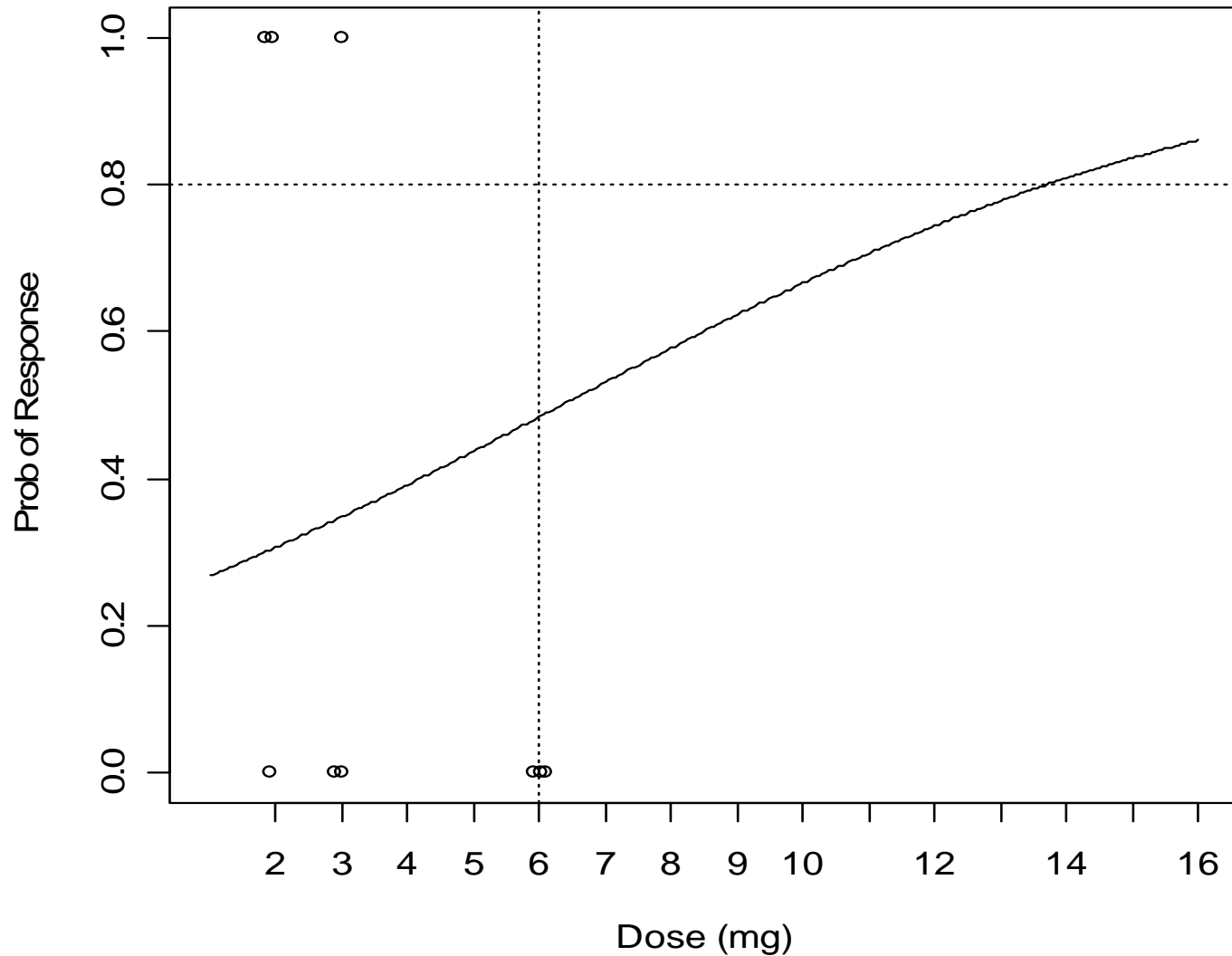


# Efficacy Example: Rapamycin in Pancreatic Cancer

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- Outcome: response
- Response = 80% inhibition of pharmacodynamic marker
- Assumption: as dose increases, % of patients with response will increase
- Desired proportion responding: 80%

# Efficacy Example: Rapamycin in Pancreatic Cancer



# Safety and Efficacy

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- Zhang, Sargent, Mandrekar
- Example: high dose can induce “over-stimulation”
- Three categories:
  - 1 = no response, no DLT
  - 2 = response, no DLT
  - 3 = DLT
- Use the continuation ratio model
- Very beautiful(!)
- Not particularly friendly at the current time for implementation

# Safety and Efficacy

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- I'm working on less beautiful, more practical approach
- $Y = 1$  if toxicity  
= 0 if no toxicity
- $Z = 1$  if efficacy  
= 0 if no efficacy
- Simultaneously search for doses with constraints based on toxicity



# Summary: “Novel” Phase I trials

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- Offer significant improvements over “traditional” phase I design
  - Safer
  - More accurate
- Slightly larger phase I: worth it!
- Related methods: Bayesian Adaptive



# Why isn't everyone using these?

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- Change in paradigm
- Larger N
- “I just want a quick phase I”
- Large investment of time from statistician
- Need time to “think” and plan it.
- IRB and others (e.g. CTEP) worry about safety (unjustified!)

# References

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