

Dose-Finding Clinical Trials in Stan

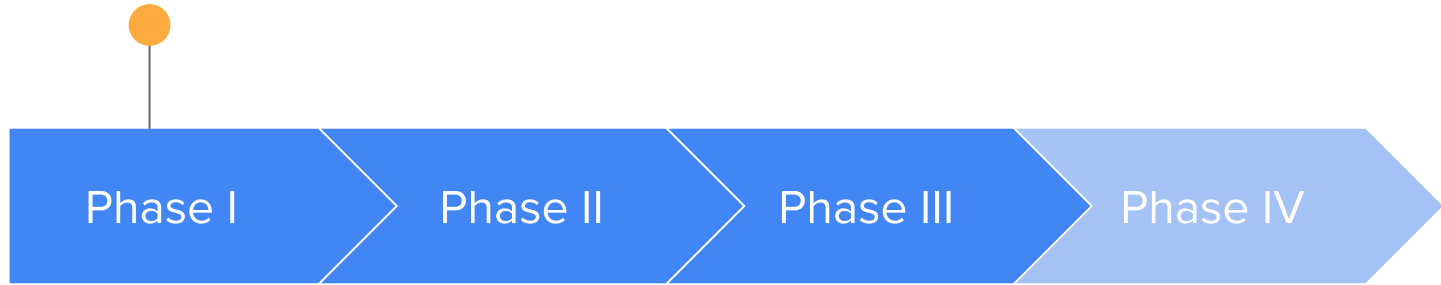
Kristian Brock, StanCon Helsinki, 30-Aug-18



What dose?

Not too toxic...but

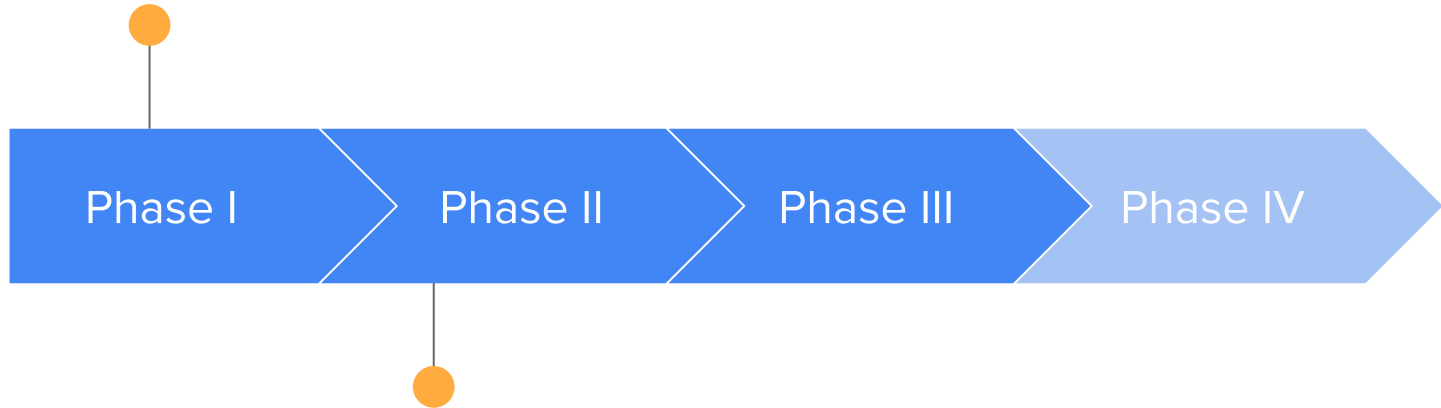
High enough to be effective



What dose?

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Is it any good?

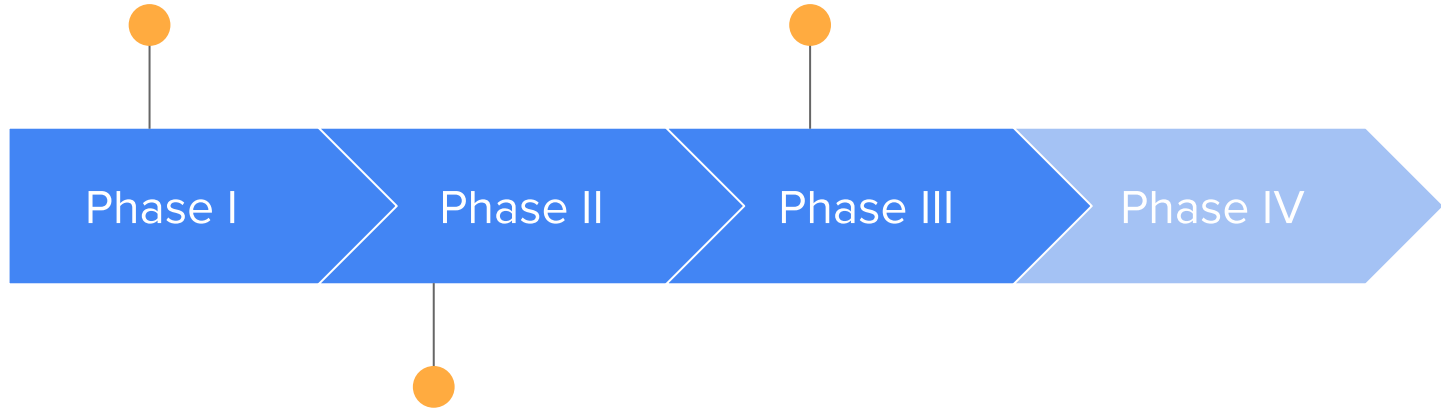
Does it do something desirable?

What dose?

Not too toxic...but
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Pivotal study

The challenger vs the standard of care



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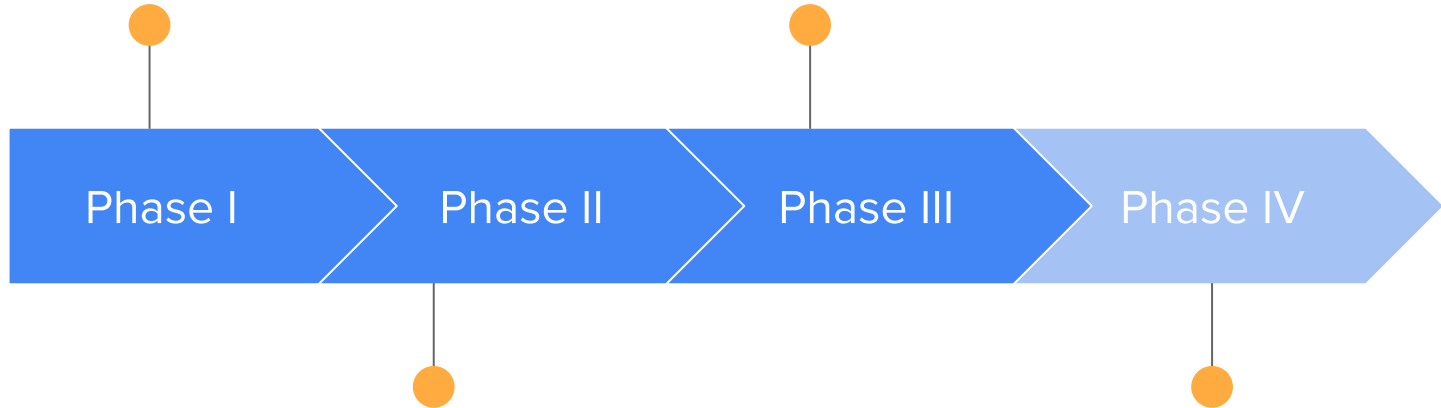
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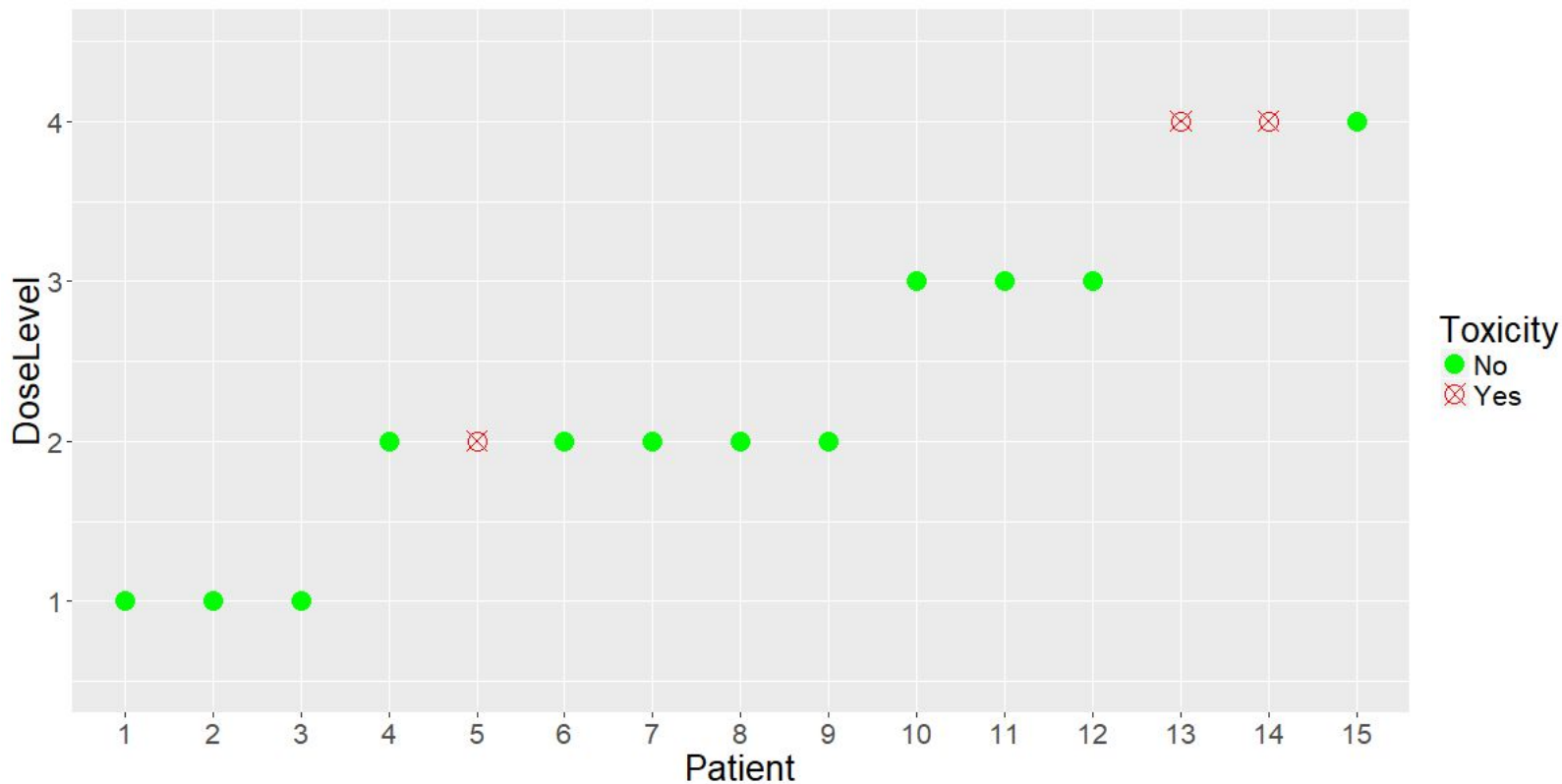
Does it do something desirable?

Post-marketing

We'll use it...
But let's keep an eye on it

**Dose-finding by
toxicity**

Dose Finding and the 3+3 Method



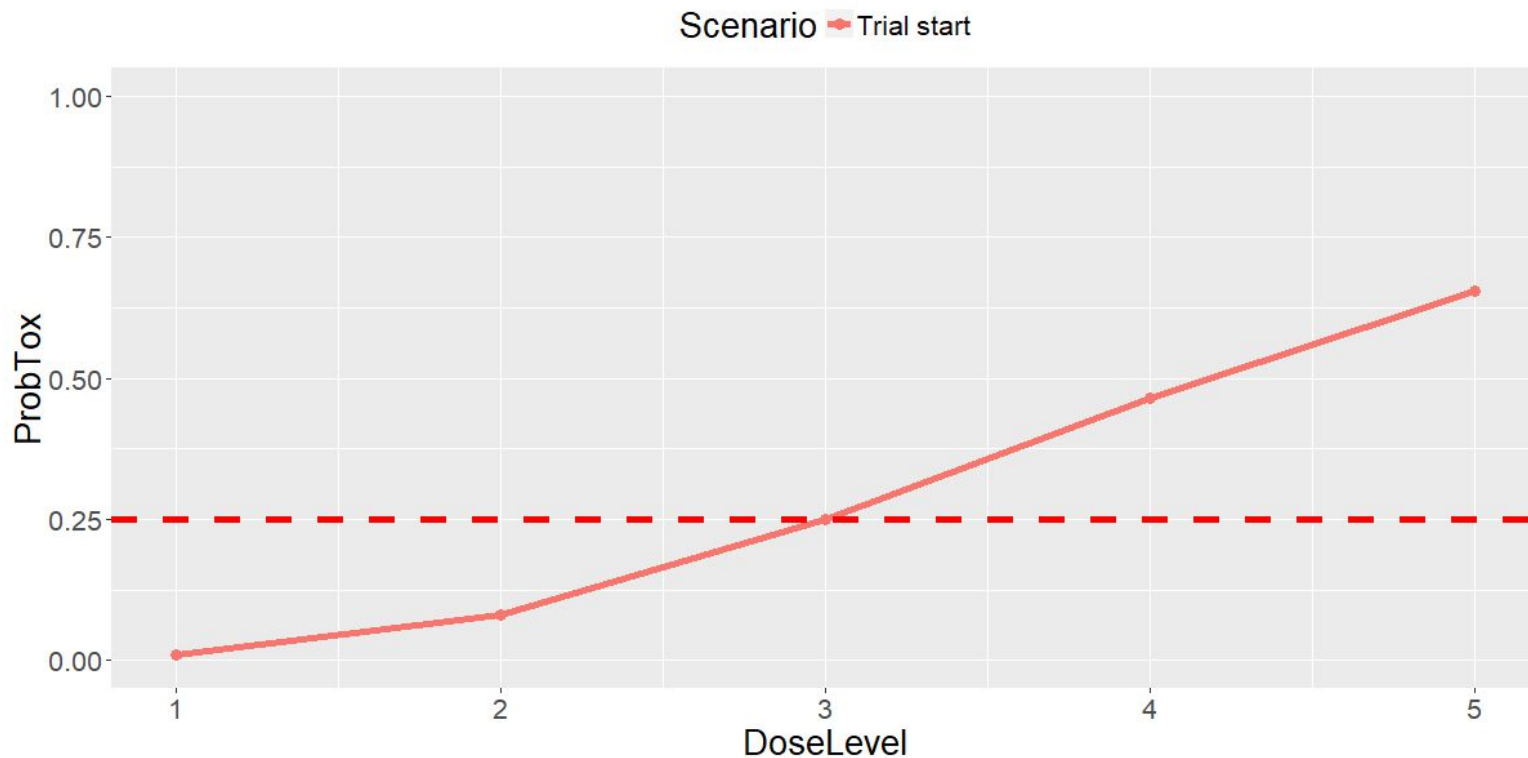
Surely folk use statistical models these days?

- Chiuzan *et al.* (2017) conducted systematic review of dose-finding methods
- Cancer trials published between 2008 and 2014

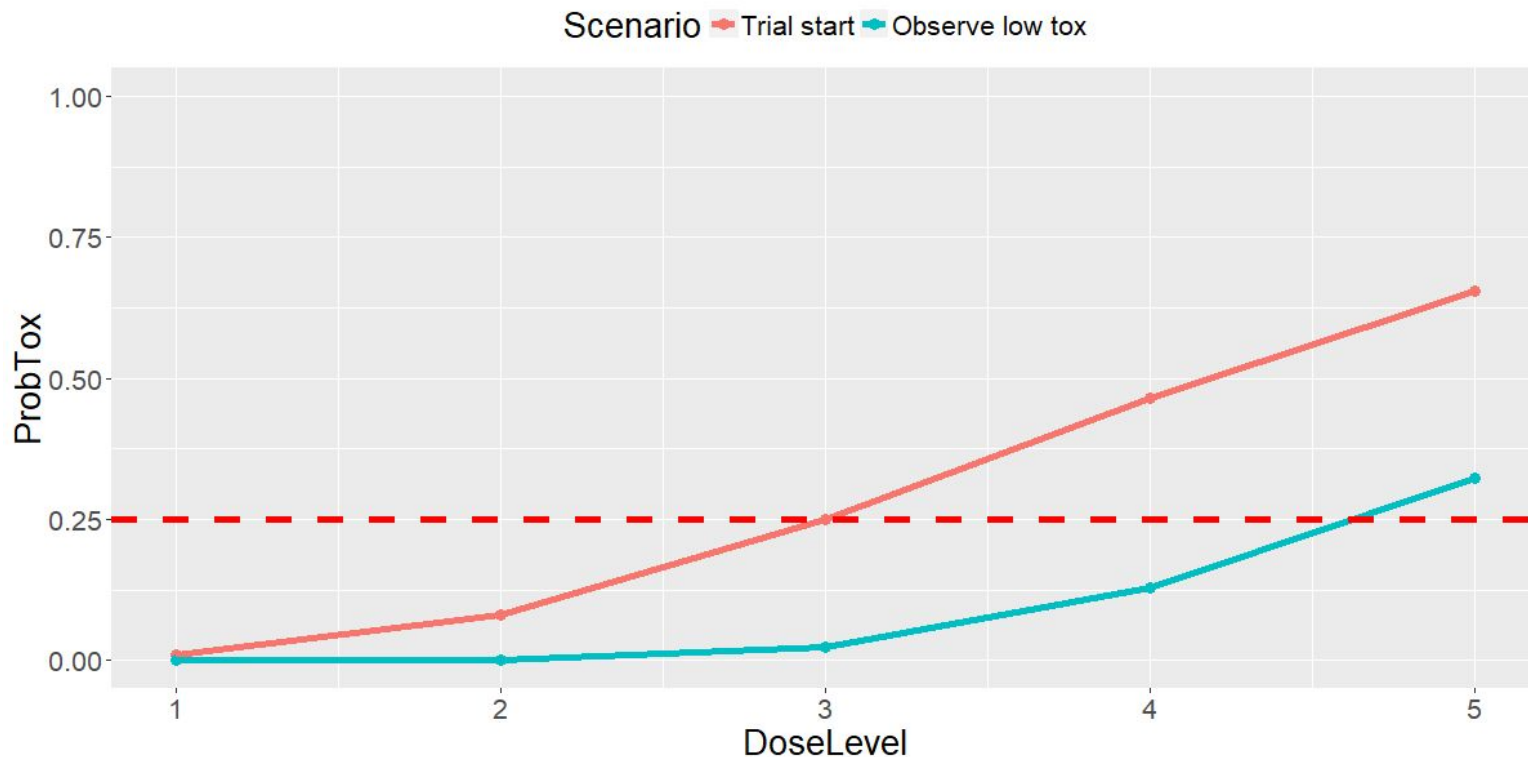


“adoption of [model-based] designs continues to remain low”

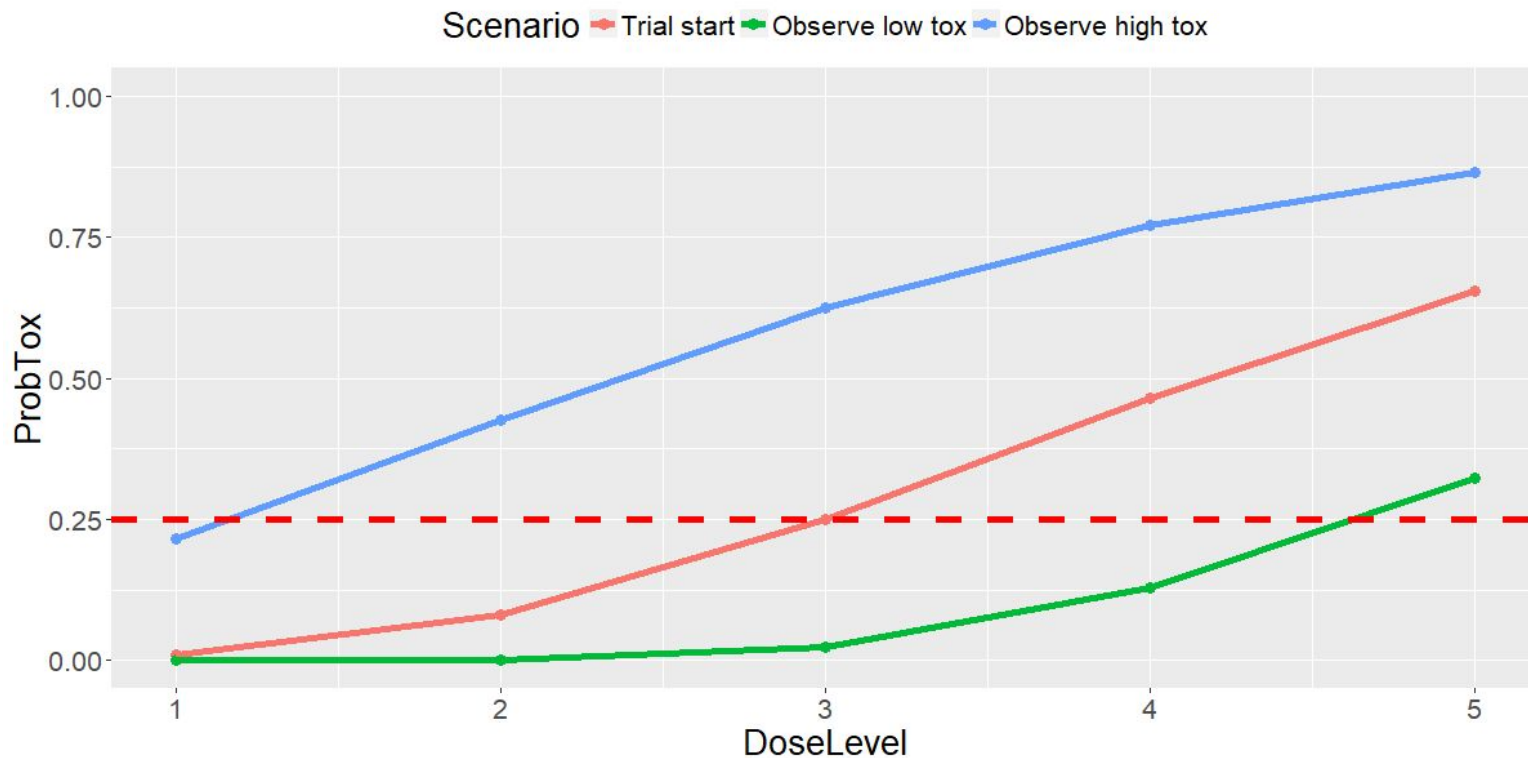
Continual Reassessment Method (CRM)



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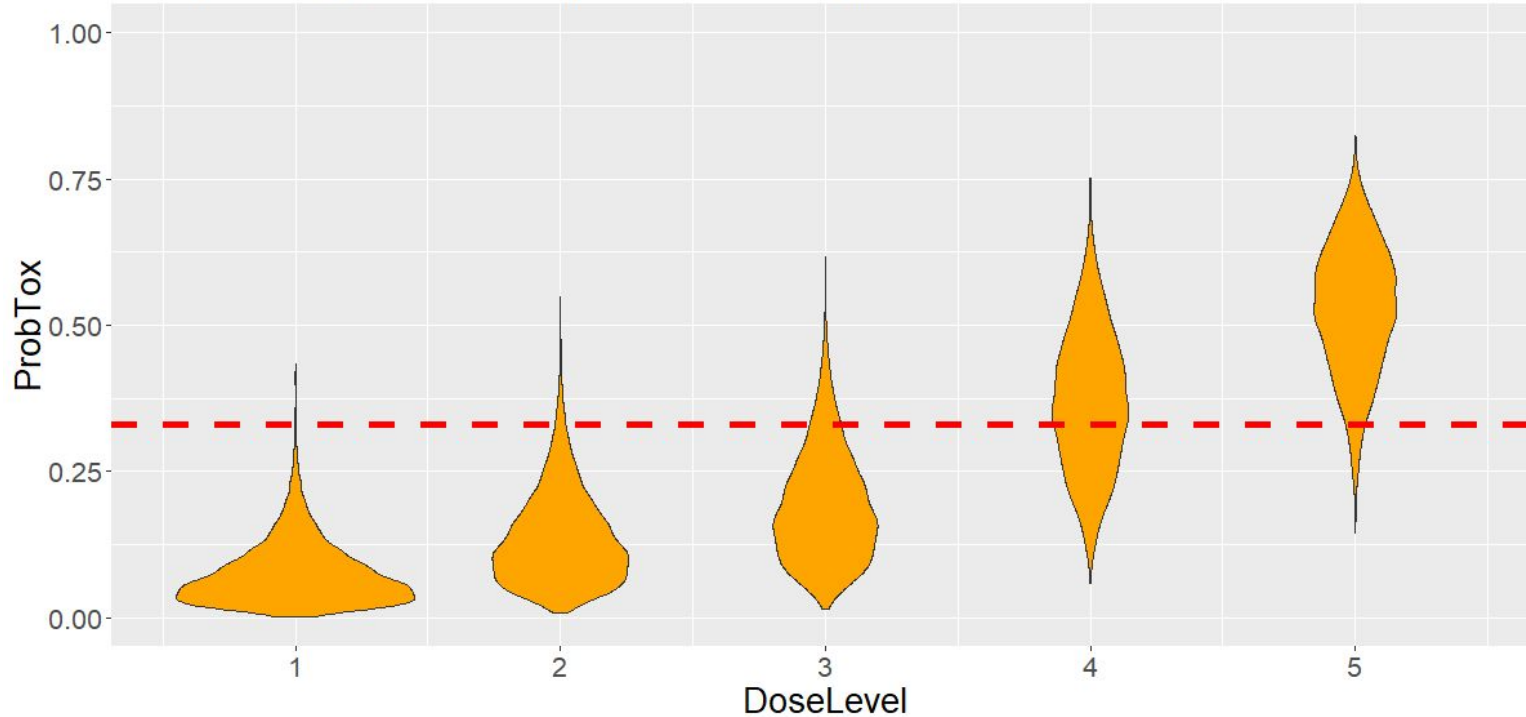


Continual Reassessment Method (CRM)



Levy et al. - ssHHT in leukaemia

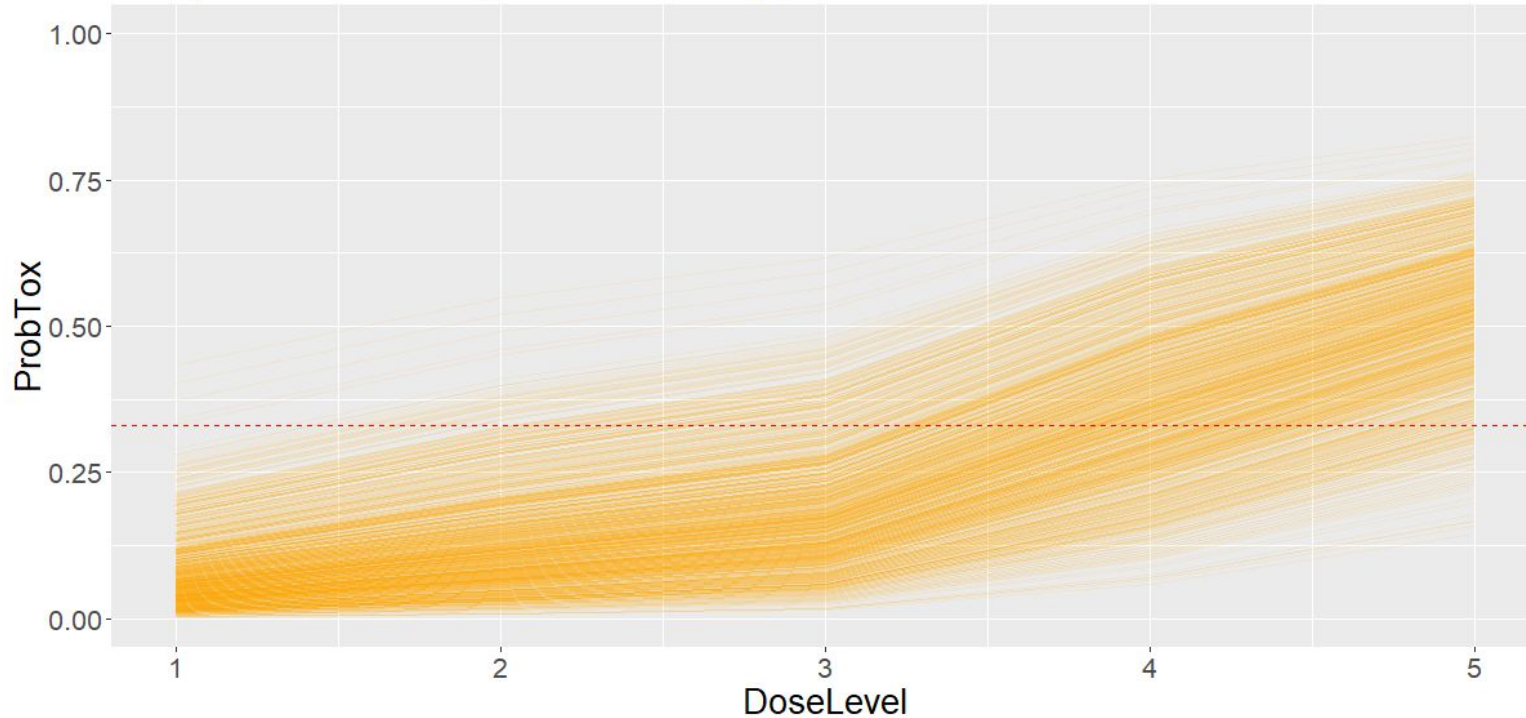
Prob(Toxicity) after all 18 patients in Levy, et al.



Levy, et al. A Phase I Dose-Finding and Pharmacokinetic Study of Subcutaneous Semisynthetic Homoharringtonine (ssHHT) in Patients with Advanced Acute Myeloid Leukaemia. *BJC*. <https://doi.org/10.1038/sj.bjc.6603265>.

Levy et al. - ssHHT in leukaemia

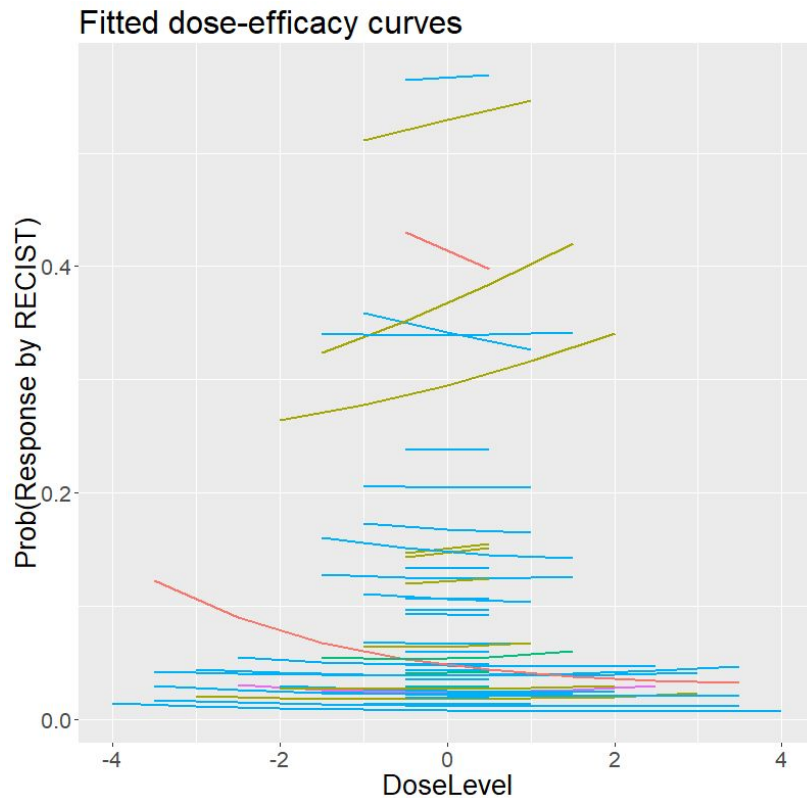
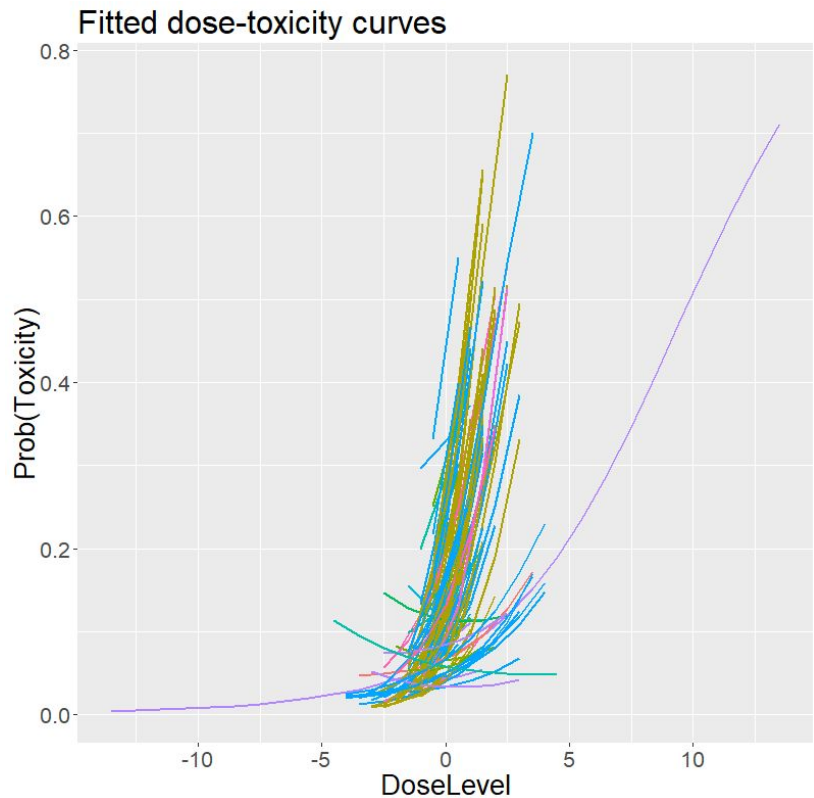
Sampled dose-toxicity curves in Levy, et al.



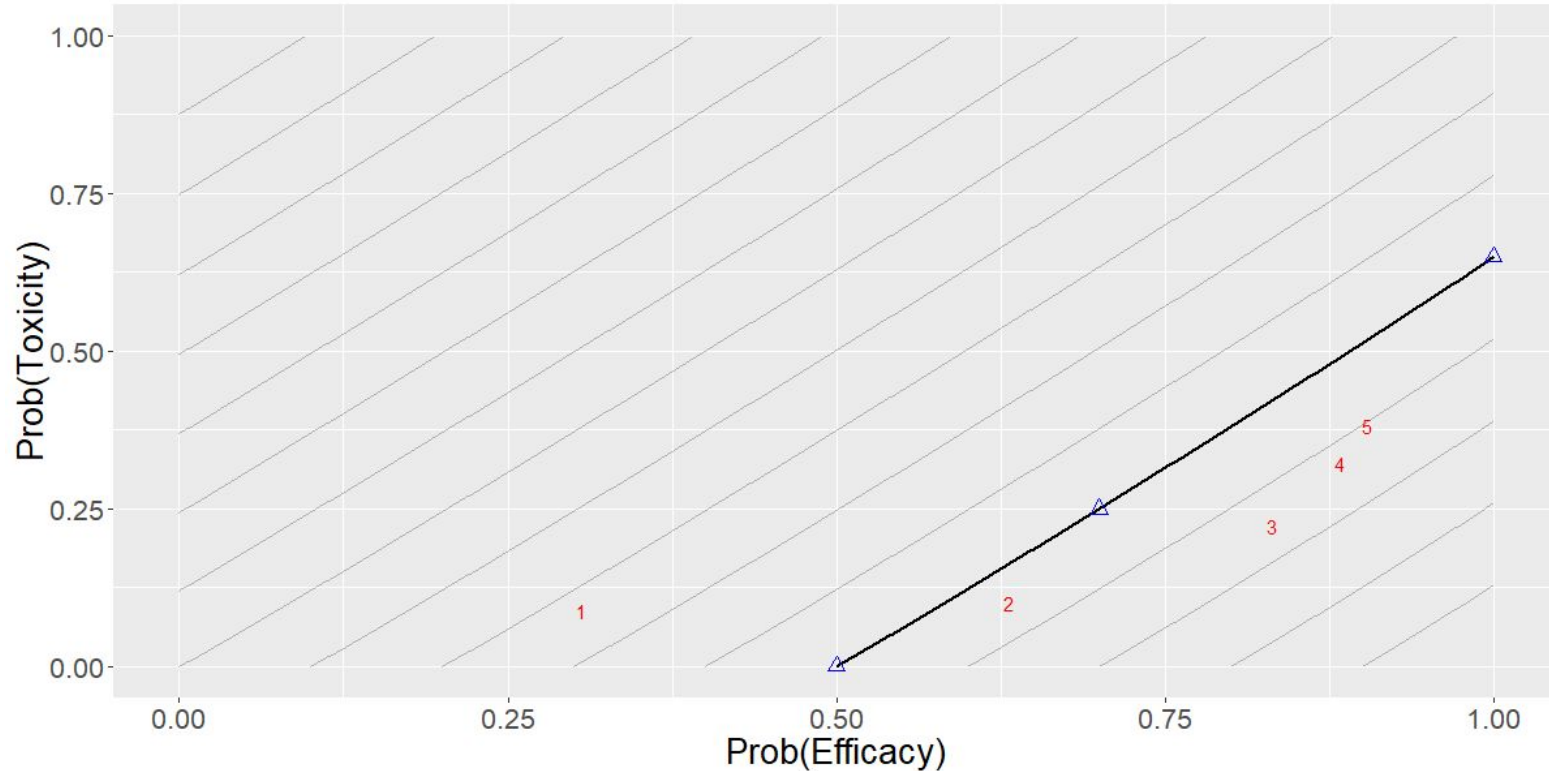
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**Dose-finding by
toxicity & efficacy**

Dose-toxicity & efficacy in modern therapies



EffTox - dose-finding based on efficacy & toxicity trade-offs



Posterior beliefs on attractiveness

Prob(“dose in column is more attractive than dose in row”)

Dose	1	2	3	4	5
1		0.92	0.86	0.82	0.8
2			0.75	0.68	0.64
3				0.58	0.54
4					0.52
5					

trialr - clinical trial designs in R & Stan

```
1
2 # Install|
3 devtools::install_github('brockk/trialr')
4 # Or
5 install.packages('trialr')
6
7 # CRM
8 levy <- '1NNN 3NNT 4NNT 4NNN 4NTN 4TNT'
9 skeleton <- c(0.05, 0.10, 0.15, 0.33, 0.5)
10 target <- 0.33
11 levy_mod <- stan_crm(levy, skeleton = skeleton, target = target,
12                    model = 'logistic_gamma', a0 = 4,
13                    beta_shape = 1, beta_inverse_scale = 1,
14                    seed = 123, control = list(adapt_delta = 0.95))
15
16 # EffTox
17 outcomes <- '1NEN 2NBE'
18 mod2 <- stan_efftox_demo(outcomes, seed = 123)
19 # That is short-hand for
20 mod2 <- stan_efftox(outcomes,
21                    real_doses = c(1.0, 2.0, 4.0, 6.6, 10.0),
22                    efficacy_hurdle = 0.5, toxicity_hurdle = 0.3,
23                    p_e = 0.1, p_t = 0.1,
24                    eff0 = 0.5, tox1 = 0.65,
25                    eff_star = 0.7, tox_star = 0.25,
26                    alpha_mean = -7.9593, alpha_sd = 3.5487,
27                    beta_mean = 1.5482, beta_sd = 3.5018,
28                    gamma_mean = 0.7367, gamma_sd = 2.5423,
29                    zeta_mean = 3.4181, zeta_sd = 2.4406,
30                    eta_mean = 0, eta_sd = 0.2,
31                    psi_mean = 0, psi_sd = 1,
32                    seed = 123)
33
```

Future work:

- Add more trial designs :-)
- Plumb it to work with tidybayes
- More visualisation via bayesplot
- Scrutinise fit by shinystan, etc
- Automated documents via Flexdashboards?

The End

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