



MRC
Biostatistics
Unit



UNIVERSITY OF
CAMBRIDGE

Adaptive Methods in Clinical Research

Lecture 1: Single arm binary outcome designs

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Ben Goldacre, Guardian 1-09-08:

- **Before 1935:** doctors were basically useless

Ben Goldacre, Guardian 1-09-08:

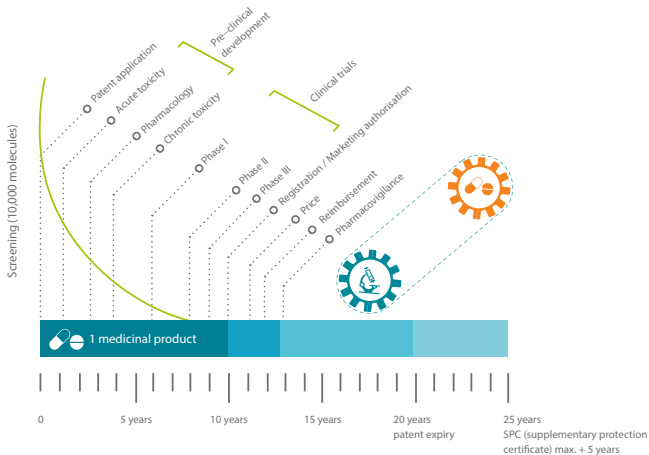
- **Before 1935:** doctors were basically useless
- **1935-1995:** antibiotics, dialysis, transplants, intensive-care units, heart surgery, every drug you've ever heard of

Ben Goldacre, Guardian 1-09-08:

- **Before 1935:** doctors were basically useless
- **1935-1995:** antibiotics, dialysis, transplants, intensive-care units, heart surgery, every drug you've ever heard of
- **1995-now:** the low-hanging fruit of medical research has all been harvested, and the industry is rapidly running out of new drugs

The development process

PHASES OF THE RESEARCH AND DEVELOPMENT PROCESS



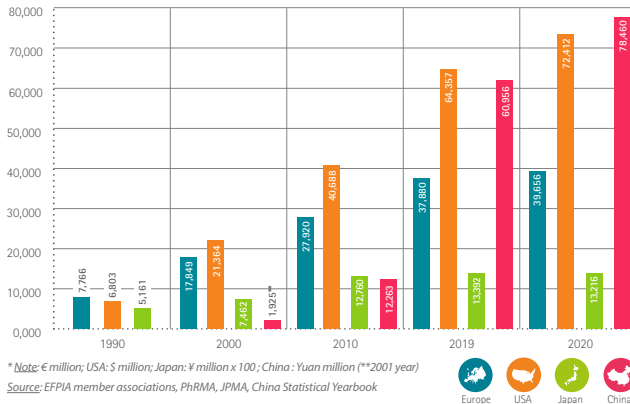
Source: European Federation of Pharmaceutical Industries and Associations (2022)

Development of a novel medicinal product

- takes 10-15 years
- costs several hundred million euros on average
 - ▶ largest contributors are confirmatory (phase III) trials
 - ▶ often involve thousands of patients with follow-up period frequently lasting years

Cost on R&D in Pharmaceutical industry

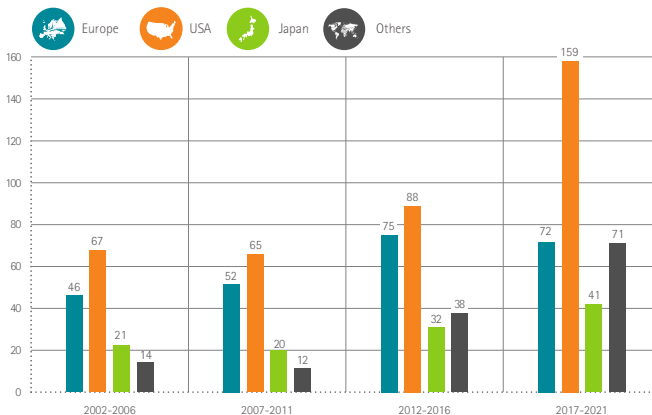
PHARMACEUTICAL R&D EXPENDITURE IN EUROPE, USA, JAPAN AND CHINA
(MILLION OF NATIONAL CURRENCY UNITS)*, 1990-2020



Source: European Federation of Pharmaceutical Industries and Associations (2022)

New molecular entities

NUMBER OF NEW CHEMICAL AND BIOLOGICAL ENTITIES (2002-2021)

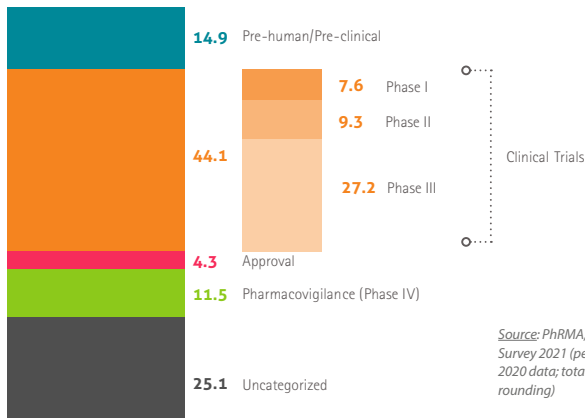


Source: SCRIIP – EFPIA calculations (according to nationality of mother company)

Source: European Federation of Pharmaceutical Industries and Associations (2022)

Cost on R&D in Pharmaceutical industry

ALLOCATION OF R&D INVESTMENTS BY FUNCTION (%)



Source: PhRMA, Annual Membership Survey 2021 (percentages calculated from 2020 data; total values may be affected by rounding)

Source: European Federation of Pharmaceutical Industries and Associations (2022)

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.

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have been unsuccessful.

- **13.4%** of treatments entering Phase I receive approval
- In oncology only **3.4%** of treatments entering Phase I receive approval

- Avoid going straight into large and expensive phase III trials
- Take more care during phases I and II
- Explore the potential of “new” statistical methods:
 - ▶ Sequential designs
 - ▶ Adaptive designs
 - ▶ Bayesian methods

What is an adaptive design?

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Possible adaptations:

- Stop the trial early when we observe a very strong (or weak) treatment effect;
- Re-estimate the required sample size;
- Stop recruitment to a poorly-performing treatment.

Why consider adaptive designs?

Different benefits for different forms of adaptation, including:

- Stopping early: fewer participants required on average
- Sample size re-estimation: more likely to reach required power
- Drop “loser” treatment: decrease the proportion of participants receiving poorly-performing treatment

In clinical trials, data accumulates steadily over time → natural to sequentially monitor results and perform interim analyses, for a number of reasons:

- **Ethical:** Minimise participants exposure to unsafe/ineffective treatments
- **Economic:** Allow *early stopping* → fewer patients needed on average.
- **Administrative:** Ensure trial is being run as planned.

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Continuous monitoring is often impractical → examine the data at periodic intervals.

Single-arm binary outcome trials

Single-arm binary outcome trials

In oncology, a single-arm binary outcome trial (response/no response) often takes place after dose finding (Phase I).

Research question: Is the response rate p of the selected dose large enough to continue development?

Single-arm binary outcome trials

We test $H_0 : p \leq p_0$ against $H_1 : p > p_0$. In general:

Single-arm binary outcome trials

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Type-I error-rate: $P(\text{reject } H_0 | H_0 \text{ is true})$

Power: $P(\text{reject } H_0 | H_1 \text{ is true})$

We wish to control the type-I error-rate to be α when $p = p_0$ and power our trial to a level $1 - \beta$ under $p = p_1$.

p_0 : the greatest response rate that we deem typical for standard of care.

p_1 : the smallest response rate that is large enough to warrant further study.

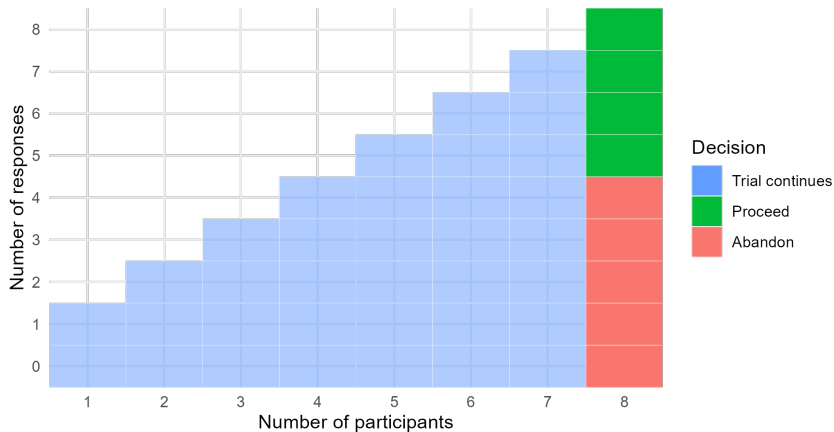
Single-stage design (A'Hern 2001)

The most simple single-arm binary outcome design is the single-stage design:

Recruit n_{max} participants. A go decision is made (i.e. the trial is deemed a success) if the final number of responses $S_{n_{max}}$ exceeds a specified boundary r (i.e. $S_{n_{max}} > r$).

We choose a set of design parameters that satisfy specified type-I error-rate and power requirements for p_0 and p_1 .

Single-stage design (A'Hern 2001)



Simon design (Simon 1989)

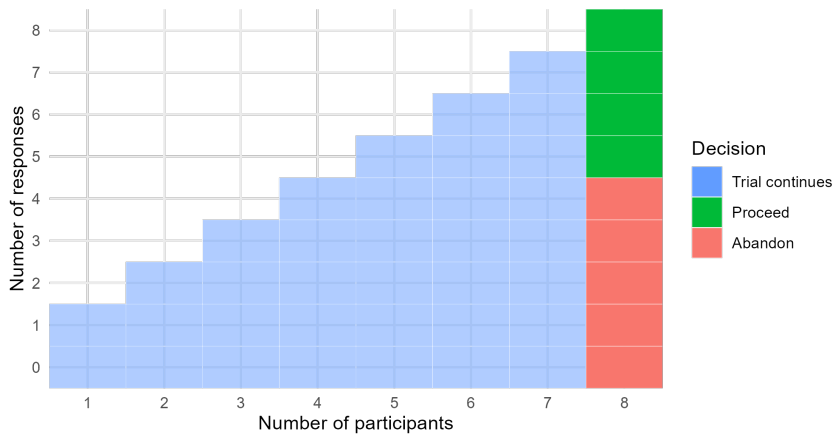
Simon design is a simple adaptation to the single-stage design:

Include single interim analysis after n_1 participants.

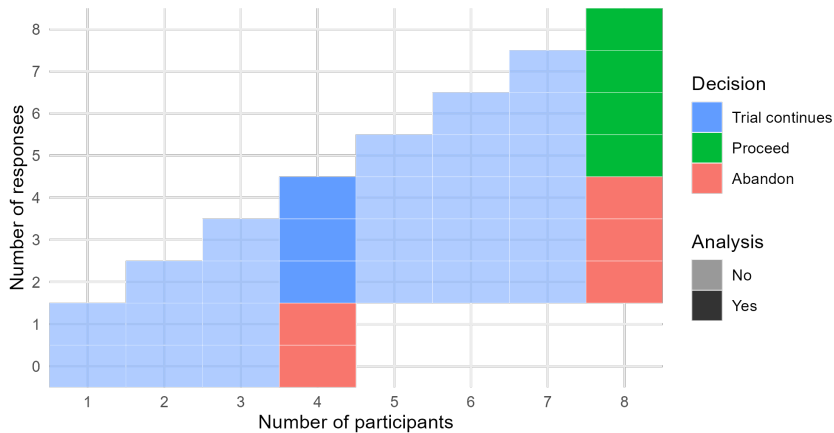
If $S_{n_1} \leq r_1$, stop for lack of benefit.

Otherwise, recruit another $n_{max} - n_1$ participants and make a go/no-go (success/failure) decision at n_{max} .

Single-stage design



Simon design



Simon designs can be obtained using the command `ph2simon` in the `clinfun` package. The command will return at least two designs:

- Minimax, which minimises the maximum sample size
- Optimal, which minimises the expected sample size under the null hypothesis ($ESS(H_0)$).

Simon design

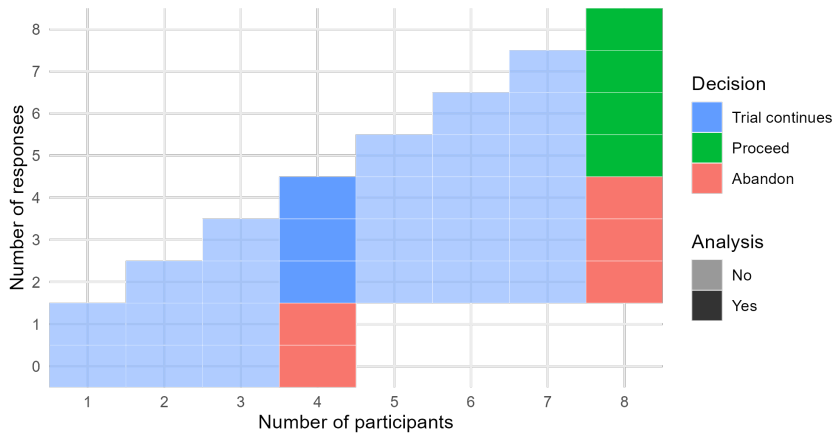
```
##
## Simon 2-stage Phase II design
##
## Unacceptable response rate: 0.2
## Desirable response rate: 0.4
## Error rates: alpha = 0.05 ; beta = 0.1
##
##           r1 n1  r  n EN(p0) PET(p0)    qLo    qHi
## Minimax      5 24 13 45  31.23  0.6559 0.108 1.000
## Admissible   4 20 14 49  30.74  0.6296 0.058 0.108
## Optimal      4 19 15 54  30.43  0.6733 0.000 0.058
```

$PET(p_0)$: Probability of Early Termination

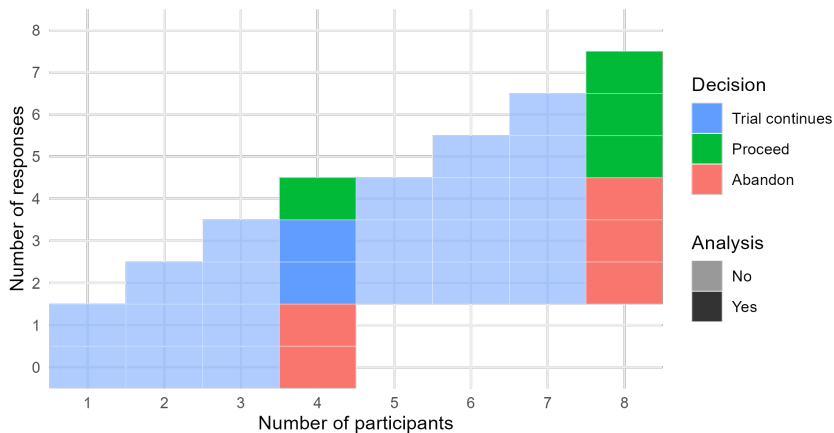
Mander and Thompson design (2010)

An extension to the Simon design, where the trial may additionally end early for a go decision after n_1 participants if $S_{n_1} \geq e_1$.

Simon design



Mander and Thompson design



Obtaining design parameters

For these binary outcome designs, it is possible to calculate the probability of rejecting H_0 using the set of design parameters, e.g. $\{n_1, n_{max}, r_1, r\}$, conditional on a response rate.

As such, we do not directly calculate stopping boundaries. Instead, we can calculate type-I error-rate as $P(\text{reject } H_0 | p = p_0)$ and power as $P(\text{reject } H_0 | p = p_1)$.

With this in mind, we can calculate α and $1 - \beta$ for all possible combinations of $\{n_1, n_{max}, r_1, r\}$ to obtain a suitable set of design parameters for our choice of p_0, p_1, α and $1 - \beta$.

Curtailment in trial design

Non-stochastic curtailment

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Stop when either success or failure is *certain*.

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Stochastic curtailment

Curtailment in trial design

Non-stochastic curtailment

Stop when either success or failure is *certain*.

Stochastic curtailment

Stop when either success or failure is *very likely*.

“Very likely”?

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How do we define “very likely”?

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How do we define “very likely”?

By using conditional power:

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Stop if failure is certain: $CP=0$ (NSC)

Stop if success is very likely: $CP > \theta_E$ (SC)

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Stop if success is certain: $CP=1$ (NSC)

Stop if failure is certain: $CP=0$ (NSC)

Stop if success is very likely: $CP > \theta_E$ (SC)

Stop if failure is very likely: $CP < \theta_F$ (SC)

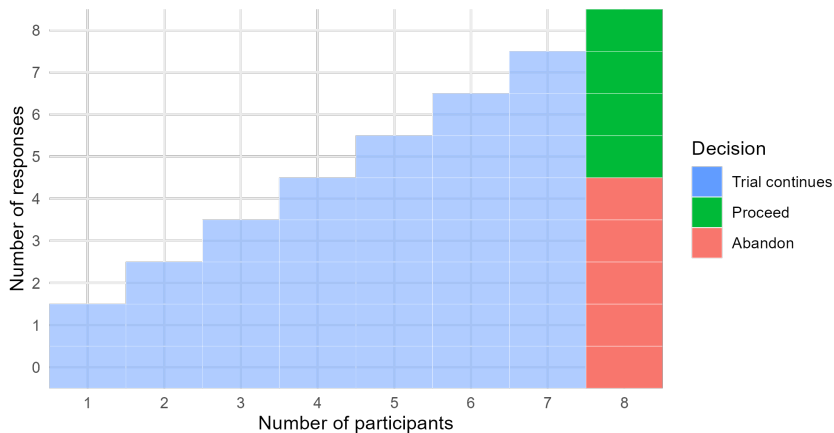
Conditional power (CP)

By planning to end a trial early due to a high or low CP, we can reduce that trial's expected sample size.

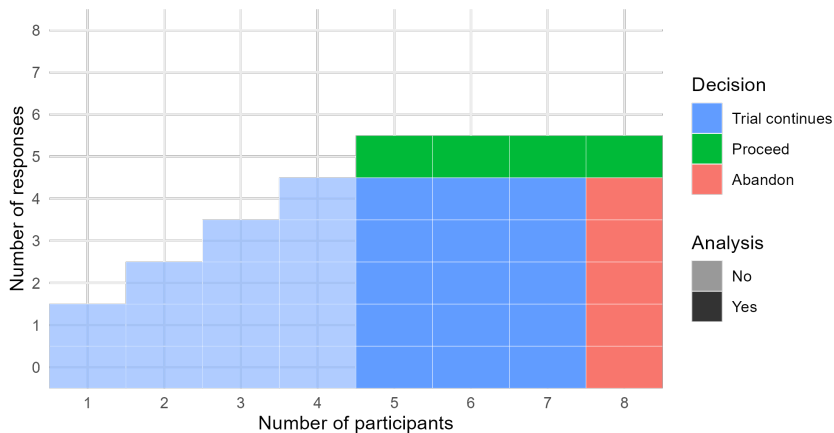
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R package for Mander and Thompson design, non-stochastic curtailment and non-stochastic curtailment designs: `curtailment`. Latest version on github (though also on CRAN), link in References.

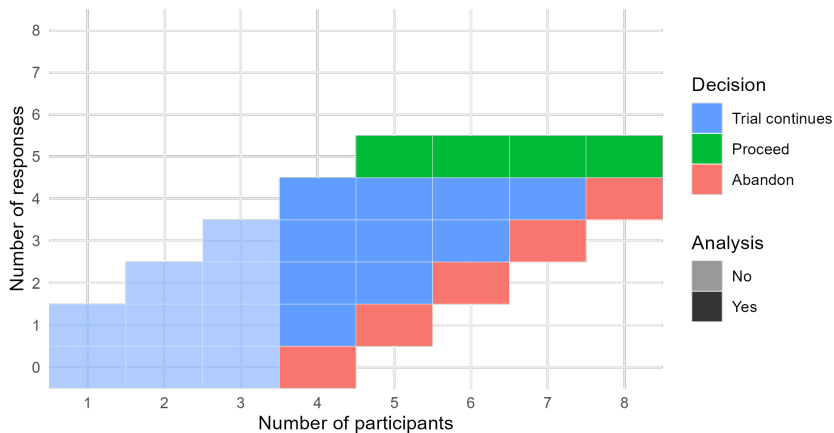
Add NSC to single-stage design



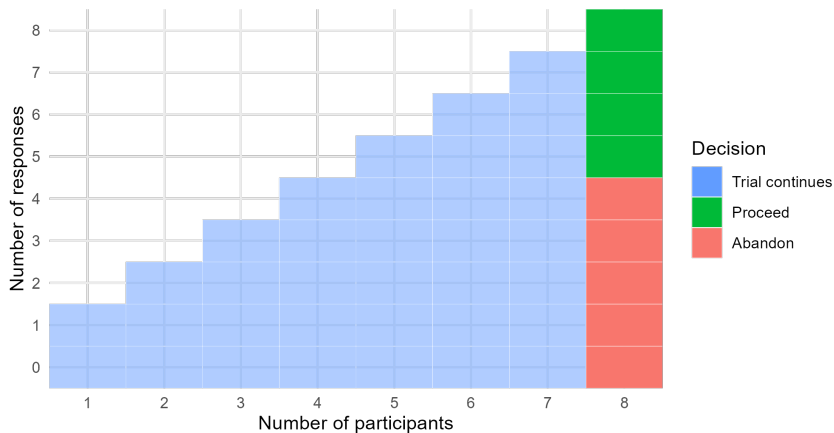
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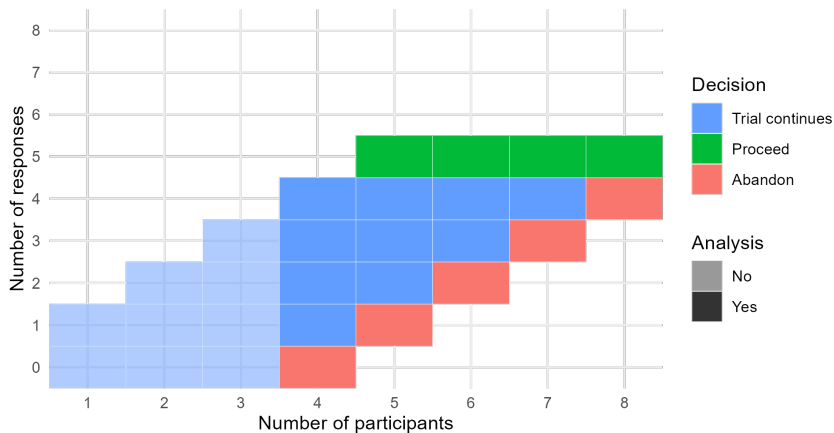
Add NSC to single-stage design



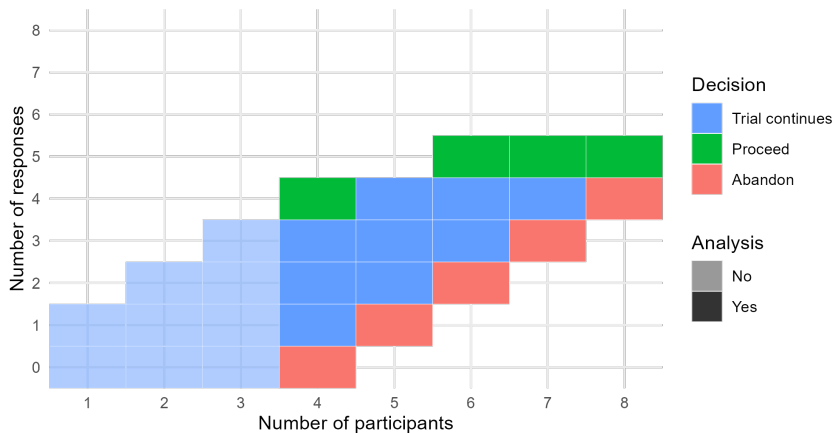
Add SC to single-stage design



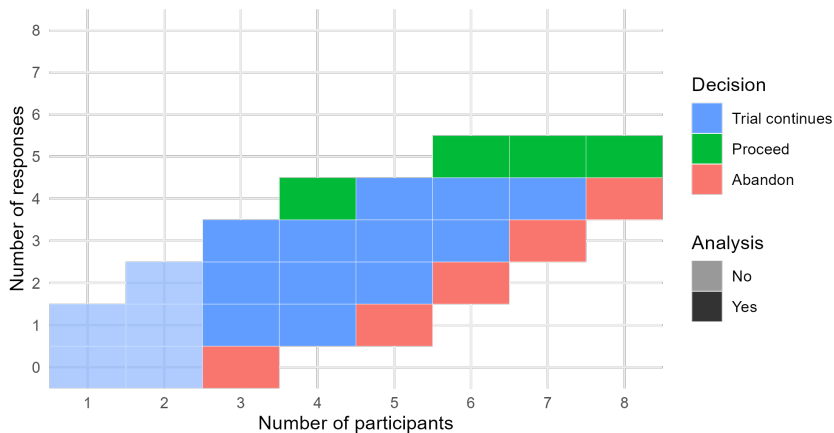
Add SC to single-stage design



Add SC to single-stage design



Add SC to single-stage design



Advantages:

- Simple, well-known to regulators
- Decreases number of participants receiving poorly-performing treatment

Disadvantages:

- When treatment works, trial will not end early (exception: you make a type II error!)
- At the points where early stopping takes place, what would be the probability of trial success if there was no stopping boundary? Not (typically) considered when choosing design parameters.

Advantages, disadvantages: Mander and Thompson

Advantages:

- Simple
- Allows early stopping for promising treatment

Disadvantages:

- May want more information if trial going well

Disadvantages:

- Final decision may be known with certainty between interim analyses
- Discrete data: may result in design with lower type-I error-rate or higher power than required.

Advantage:

- Decreased expected sample size under both $p = p_0$ and $p = p_1$.

Disadvantages:

- More frequent monitoring required:
 - ▶ More work (both analysis and logistics)
 - ▶ Difficult if responses come quickly

Beyond single-arm single-stage designs

- Curtailment can also be used in the two-arm setting
- Next lecture: multiple arms and multiple stages

References

- R.P. A'Hern. Sample size tables for exact single-stage phase II designs. *Statistics in Medicine*, 20(6):859–866, 2001
- R. Simon. Optimal two-stage designs for phase II clinical trials. *Controlled Clinical Trials*, 10:1–10, 1989.
- A.P. Mander and S.G. Thompson. Two-stage designs optimal under the alternative hypothesis for phase II cancer clinical trials. *Contemporary Clinical Trials*, 31(6):572-578, 2010.
- M. Law, M.J. Grayling, and A.P. Mander. A stochastically curtailed single-arm phase II trial design for binary outcomes” *Journal of biopharmaceutical statistics* 32.5: 671-691, 2022.
- M. Law, M.J. Grayling, and A.P. Mander. A stochastically curtailed two-arm randomised phase II trial design for binary outcomes. *Pharmaceutical Statistics*, 20.2: 212-228, 2021.
- M. Law. Github repository.