

Sample size recalculation for a skewed outcome in two-stage three-arm sequential non- inferiority clinical trials: a simulation study

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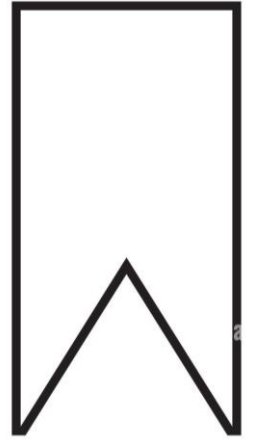


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Background



- Non-inferiority clinical trial & FDA guidelines
- The base-point of this study: the Lanyu Lei resampling (original) algorithm
- The skewed outcomes in clinical research



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Non-inferiority clinical trial & FDA guidelines (1)



Non-inferiority clinical trial & FDA guidelines (2)

Non-Inferiority Clinical Trials to Establish Effectiveness Guidance for Industry

Additional copies are available from:
Office of Communications, Division of Drug Information
Center for Drug Evaluation and Research
Food and Drug Administration
10001 New Hampshire Ave., Hillandale Bldg., 4th Floor
Silver Spring, MD 20993-0002
Phone: 855-543-3784 or 301-796-3400; Fax: 301-431-6353
Email: druginfo@fda.hhs.gov

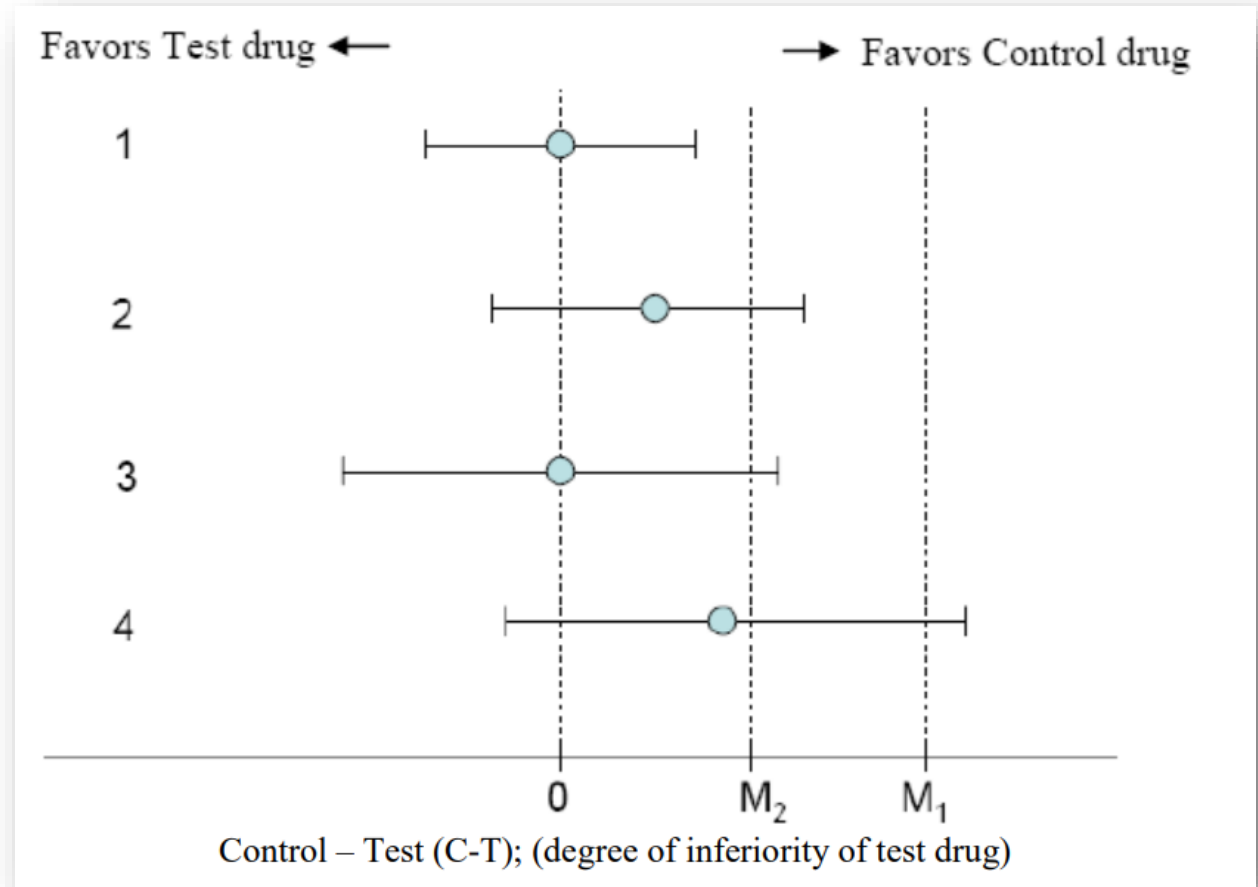
<http://www.fda.gov/Drugs/GuidanceComplianceRegulatoryInformation/Guidances/default.htm>
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Office of Communication, Outreach and Development
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10903 New Hampshire Ave., Bldg. 71, Room 3128
Silver Spring, MD 20993-0002
Phone: 800-835-4709 or 240-402-8010
Email: ocod@fda.hhs.gov

<http://www.fda.gov/BiologicsBloodVaccines/GuidanceComplianceRegulatoryInformation/Guidances/default.htm>

U.S. Department of Health and Human Services
Food and Drug Administration
Center for Drug Evaluation and Research (CDER)
Center for Biologics Evaluation and Research (CBER)

November 2016
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Non-inferiority clinical trial & FDA guidelines (3)

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- **Constancy** → The demonstrated effect of the active control over placebo in the historical trial has not changed over time
- **Assay sensitivity** → The study's ability to distinguish an effective treatment from a less effective or an ineffective one

This assumptions are not testable in a trial without a concurrent placebo group



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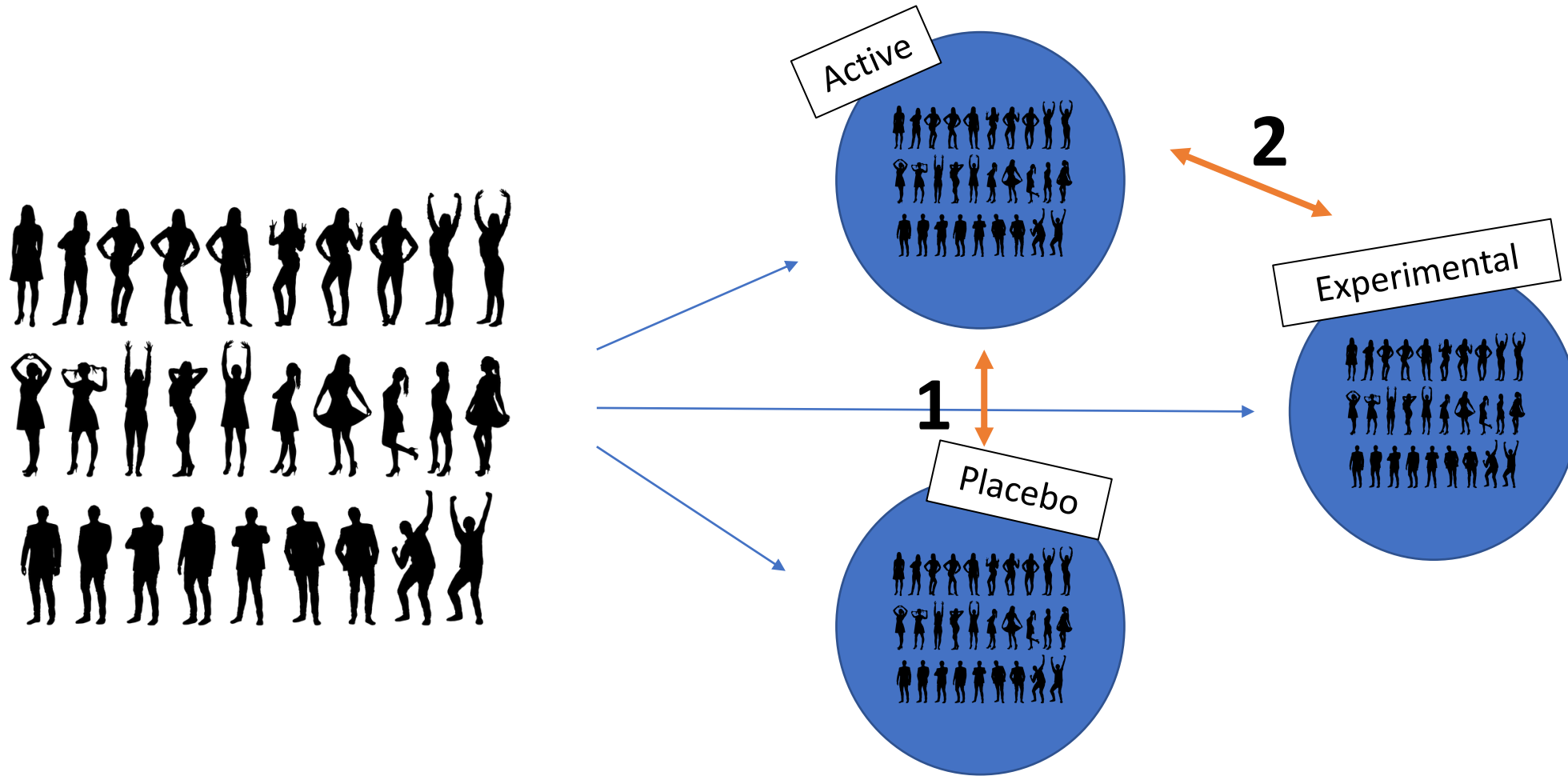


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Non-inferiority clinical trial & FDA guidelines (4)



Non-inferiority clinical trial & FDA guidelines (5)

Superiority

$$H_{0,AP}^{(s)}: \mu_A \leq \mu_P \text{ versus } H_{1,AP}^{(s)}: \mu_A > \mu_P$$

Noninferiority

Fixed Margin

or

$$H_{0,TA}^{(n)}: \mu_T \leq \mu_A - \Delta \text{ versus } H_{1,TA}^{(n)}: \mu_T > \mu_A - \Delta$$

Noninferiority

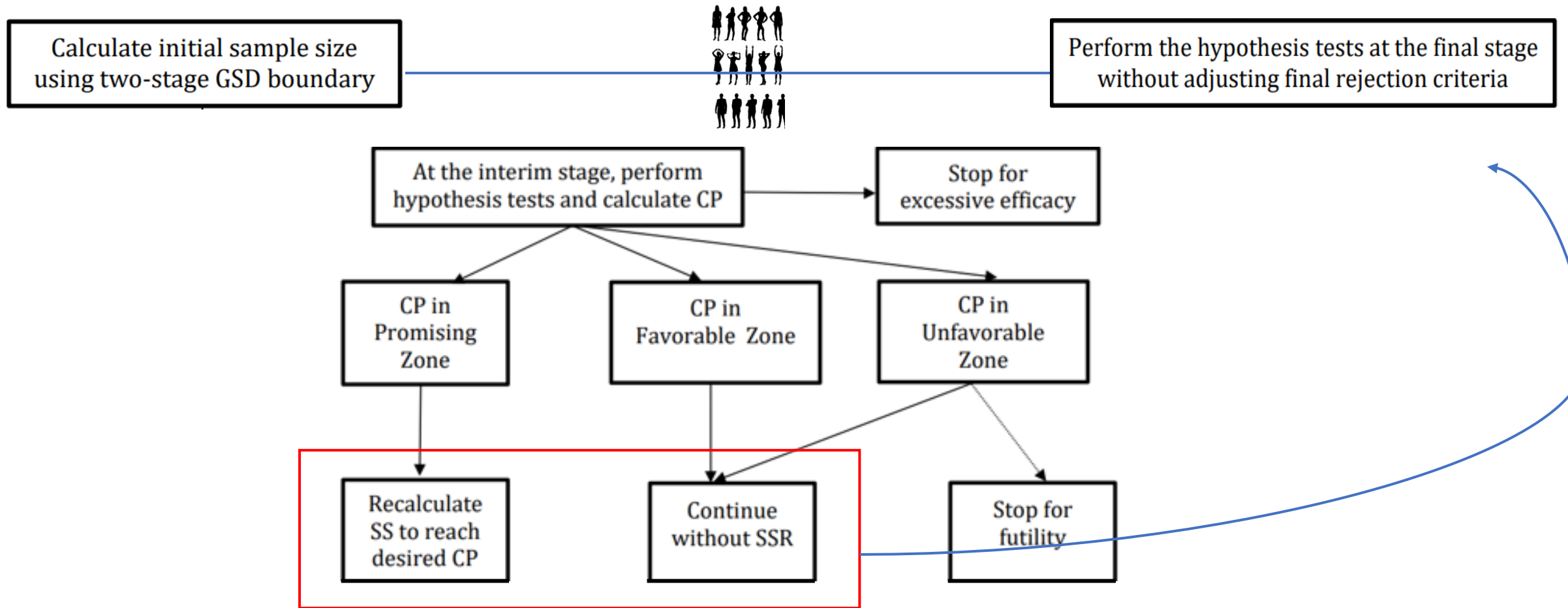
Effect of preservation

$$H_{0,TA}^{(n)}: \frac{\mu_T - \mu_P}{\mu_A - \mu_P} \leq 1 - \lambda \text{ versus } H_{1,TA}^{(n)}: \frac{\mu_T - \mu_P}{\mu_A - \mu_P} > 1 - \lambda$$

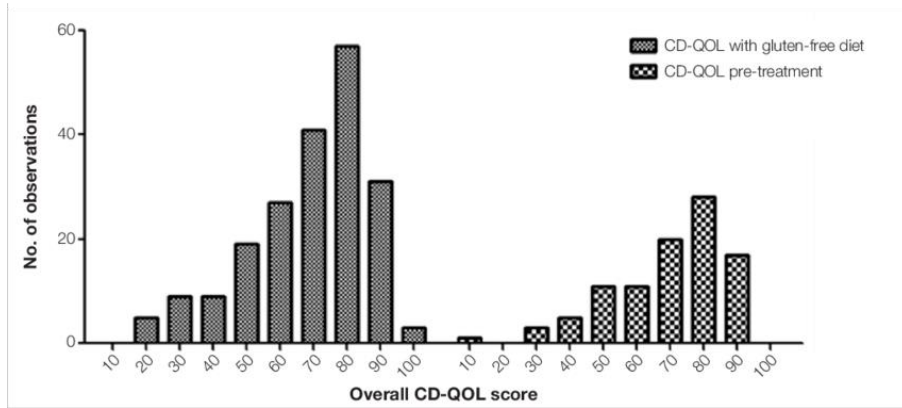


The base-point of this study: the Lanyu Lei resampling algorithm

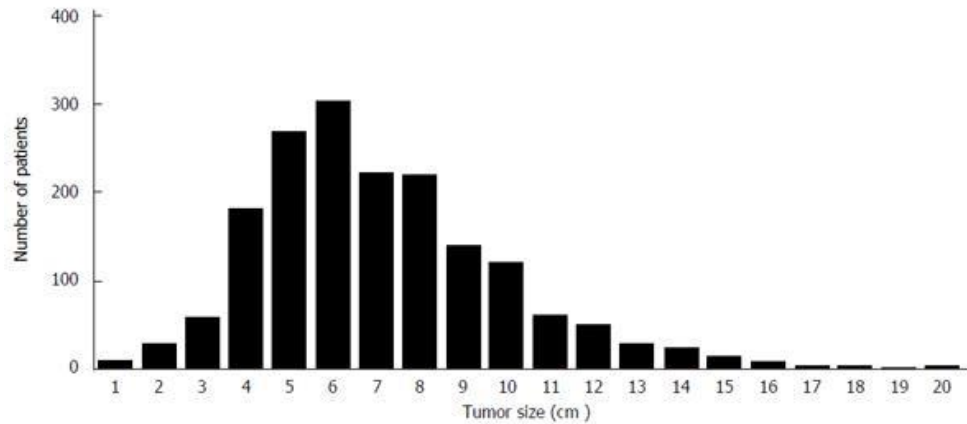
A group sequential design based on the promising zone of conditional power (1)



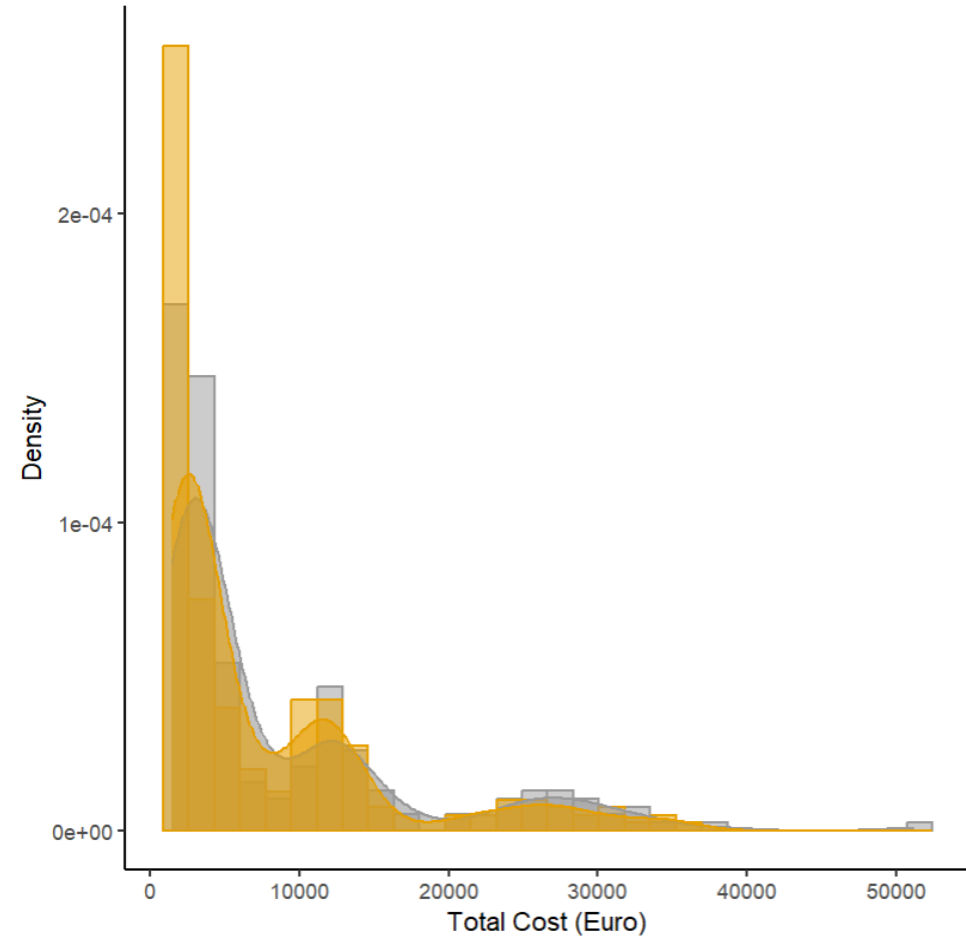
The skewed outcomes in clinical research



Casellas F. et al. (2014)



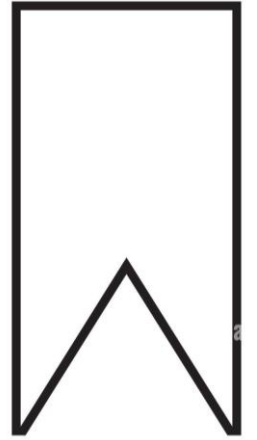
Huang B. et al. (2016)



Desideri A. et al. (2003)

Methods

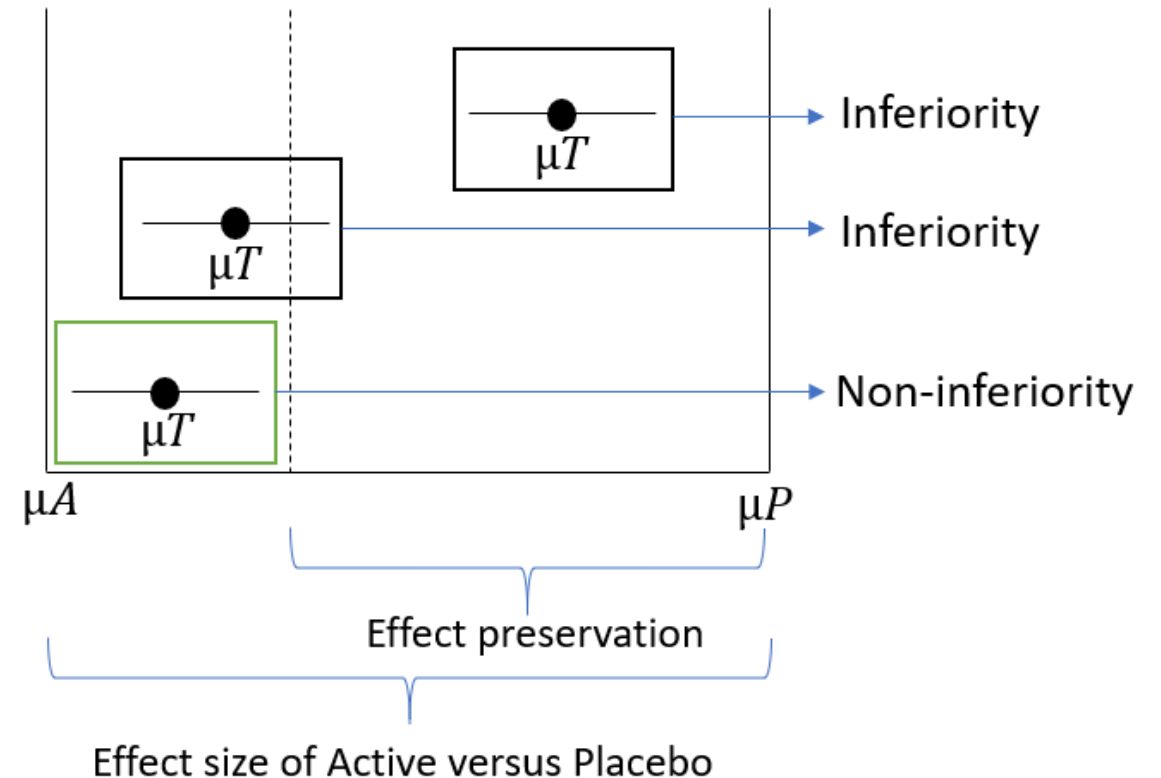
- The Coverage probability of the original sample size re-calculation method in the presence of deviation from normality
- The proposed algorithm for skewed outcomes
- A real-data example
- A brief presentation of the r-shiny web application



The coverage probability of the original sample size re-calculation method in the presence of deviation from normality

COVERAGE PROBABILITY:

The **total number of rejected nulls** over the **total number of simulations** ($N=10,000$), under the **alternative hypothesis**, that in case of “smaller is better” is:



The coverage probability of the original sample size re-calculation method in the presence of deviation from normality

Sample size



$$Y(P) \sim N(\mu_P, \sigma)$$

$$Y(A) \sim N(\mu_A, \sigma)$$

$$Y(T) \sim N(\mu_T, \sigma)$$



Simulated populations



$$Y(P) \sim \text{Gamma}(k_P, \theta_P)$$

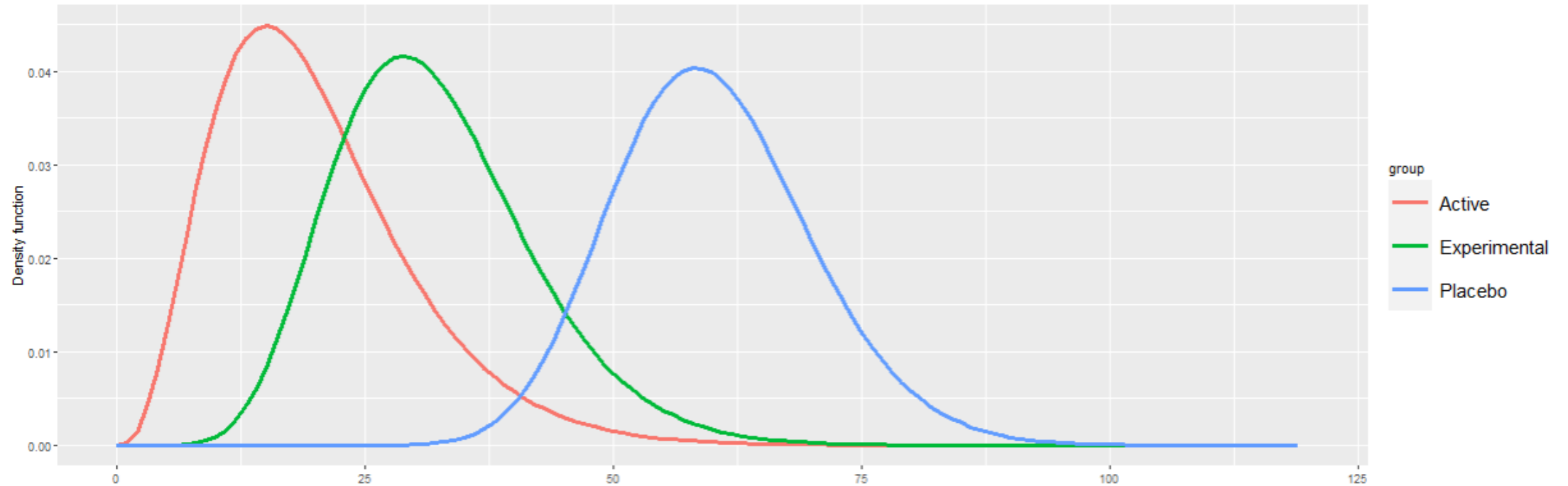
$$Y(A) \sim \text{Gamma}(k_A, \theta_A)$$

$$Y(T) \sim \text{Gamma}(k_T, \theta_T)$$

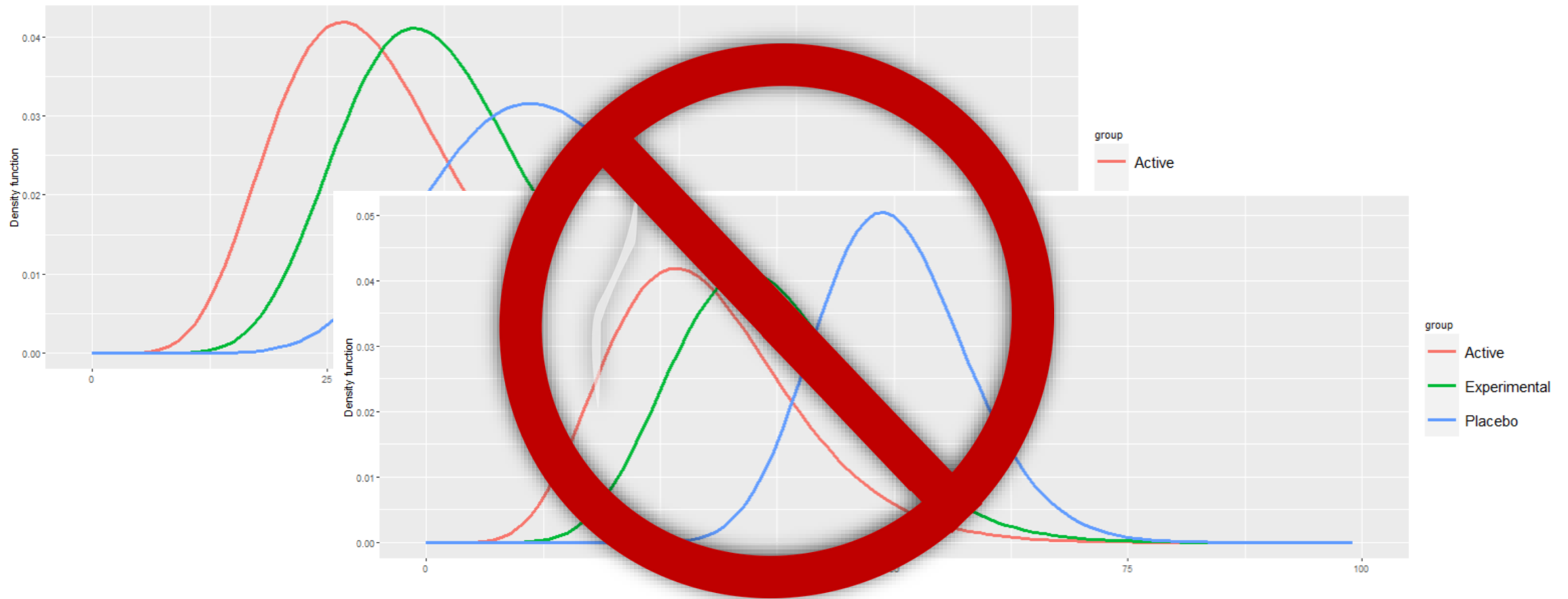
$$\begin{cases} \mu = k * \theta \\ \sigma = \sqrt{k * \theta^2} \end{cases} =$$

$$\begin{cases} k = (\mu/\sigma)^2 \\ \theta = \sigma^2/\mu \end{cases}$$

The coverage probability of the original sample size re-calculation method in the presence of deviation from normality



The coverage probability of the original sample size re-calculation method in the presence of deviation from normality



The proposed algorithm for skewed outcomes (1)

The **power function** used to estimate the initial sample size

$$1 - \beta = \phi \left(\frac{\mu T - (1 - \lambda)\mu A - \lambda\mu P}{\frac{\sigma}{\sqrt{nT}} \sqrt{1 + \frac{(1 - \lambda)^2}{cA} + \frac{\lambda^2}{cP}}} - Z_{1-\alpha(2)} \right)$$

$$1 - \beta = \phi \left(\frac{\mu T - (1 - \lambda)\mu A - \lambda\mu P}{\sqrt{\frac{Var(T)}{nT} + \frac{(1 - \lambda)^2 * Var(A)}{nT * cA} + \frac{\lambda^2 Var(P)}{nT * cP}}} - t_{1-\alpha(2),v} \right)$$



The proposed algorithm for skewed outcomes (2)

The test statistics for $H_{1,AP} = \mu_A < \mu_P$:

$$Z_{PA} = \frac{\mu_P - \mu_A}{\sigma \sqrt{\frac{1}{nT * cA} + \frac{1}{nT * cP}}}$$

$$T_{PA} = \frac{\mu_P - \mu_A}{\sqrt{\frac{Var(A)}{nT * cA} + \frac{Var(P)}{nT * cP}}}$$



The proposed algorithm for skewed outcomes (3)

- The test statistics for $H_{1, TA} = \frac{(\mu P - \mu T)}{(\mu P - \mu A)} > 1 - \lambda$:

$$Z_{TA} = \frac{(1 - \lambda)\mu A + \lambda\mu P - \mu T}{\frac{\sigma}{\sqrt{nT}} \sqrt{1 + \frac{(1 - \lambda)^2}{cA} + \frac{\lambda^2}{cP}}}$$

$$T_{TA} = \frac{(1 - \lambda)\mu A + \lambda\mu P - \mu T}{\sqrt{\frac{Var(T)}{nT} + \frac{(1 - \lambda)^2 * Var(A)}{nT * cA} + \frac{\lambda^2 Var(P)}{nT * cP}}}$$

The proposed algorithm for skewed outcomes (4)

The **conditional power** function:

$$CondP = 1 - \phi \left(\frac{Z_{1-\alpha(2)}\sqrt{nT(2)} - Z_{TA(1)}\sqrt{nT(1)}}{\sqrt{nT(2)} - nT(1)} - \frac{(1-\lambda)\mu A + \lambda\mu P - \mu T}{\frac{\sigma}{\sqrt{nT(2)} - nT(1)} \sqrt{1 + \frac{(1-\lambda)^2}{cA} + \frac{\lambda^2}{cP}}} \right)$$

$$CondP = 1 - \phi \left(\frac{t_{1-\alpha(2),v}\sqrt{nT(2)} - T_{TA(1)}\sqrt{nT(1)}}{\sqrt{nT(2)} - nT(1)} - \frac{(1-\lambda)\mu A + \lambda\mu P - \mu T}{\sqrt{\frac{Var(T)}{nT(2)} + \frac{(1-\lambda)^2 * Var(A)}{(nT(2) - nT(1)) * cA} + \frac{\lambda^2 Var(P)}{(nT(2) - nT(1)) * cP}}} \right)$$



A real data example (1)

European Heart Journal (2003) 24, 1630–1639



Cost of strategies after myocardial infarction (COSTAMI)

A multicentre, international, randomized trial for
cost-effective discharge after uncomplicated myocardial
infarction

Alessandro Desideri^{a*}, Paolo Maria Fioretti^b, Lauro Cortigiani^c,
Dario Gregori^d, Claudio Coletta^e, Carlo Vigna^f, Francesco Tota^g,
Riccardo Rambaldi^h, Jeroen Baxⁱ, Leopoldo Celegon^j, Riccardo Bigi^k,
Eugenio Picano^l, on behalf of the COSTAMI (Cost of Strategies After
Myocardial Infarction) Trial Investigators



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A real data example (2)

Groups	Total cost	
	Mean	Standard Deviation
Usual Care Strategy= Placebo/control group	8344	8894
Early Discharge=active comparator group	7391	7903

$$\mu_P - \mu_A = \text{€ } 953$$

$$1 - \lambda = 0.5$$



$$\mu_P - \mu_T > 953 * 0.5 = \text{€ } 476.5$$

$$\mu_T < 8344 - 476.5 = 7867.5$$



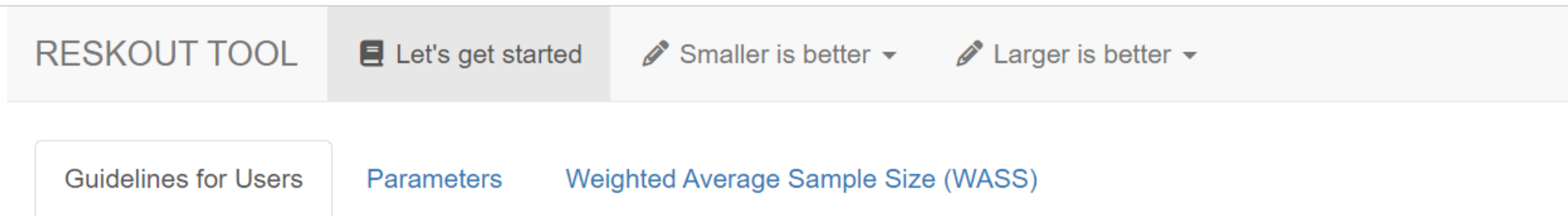
A real data example (3)

Groups	Total cost	
	Mean	Standard Deviation
Usual Care Strategy= control group	8344	8894
Early Discharge=active comparator group	7391	7903
New strategy= experimental group	7666	7488



A brief presentation of the r-shiny web application

Resampling algorithm for **SK**ewed **OUT**come



Sample size re-calculation for a skewed outcome in two-stage three-arm sequential noninferiority clinical trials: a simulation study

<https://r-ubesp.dctv.unipd.it/shiny/reskout/>



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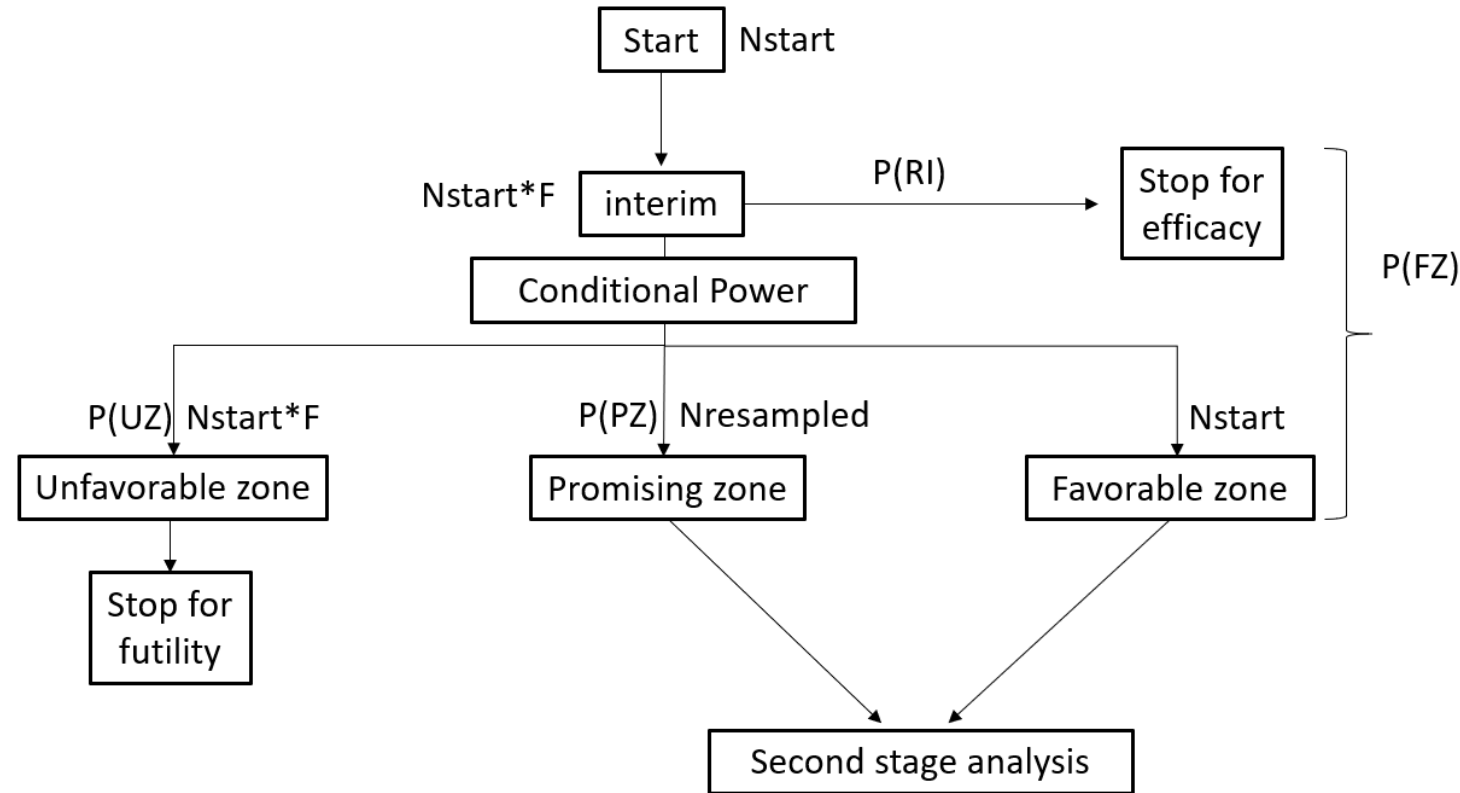


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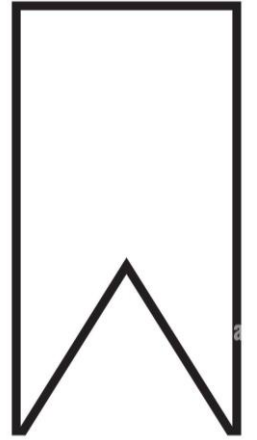
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Diagram of the resampling algorithm & the weighted average sample size (WASS)



$$(Nstart * F) * P(RI) + Nstart * (P(FZ) - P(RI)) + (Nstart * F) * P(UZ) + Nresampled * P(PZ)$$

Results



- Comparison between study designs under diverse hypothesis settings
- Study design by original and proposed algorithm by the real-data example



Comparison between study designs under diverse hypothesis (1)

Estimates under SD=SDP		Estimates under SDP>SDA by the proposed method									
		0%		10%		20%		30%		40%	
N	Nr	N	Nr	N	Nr	N	Nr	N	Nr	N	Nr
172	189	150	166	128	142	110	122	97	109	86	97
CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)
0.84	0.2	<u>0.84</u>	<u>0.17</u>	<u>0.85</u>	<u>0.16</u>	<u>0.85</u>	<u>0.16</u>	<u>0.85</u>	<u>0.14</u>	<u>0.85</u>	<u>0.13</u>



Comparison between study designs under diverse hypothesis (2)

Sample size assuming SD=SDP	CP and P(RI) assuming SD=SDP when SDA>SDP									
	10%		20%		30%		40%		50%	
	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)
172	<u>0.77</u>	<u>0.12</u>	<u>0.72</u>	<u>0.09</u>	<u>0.66</u>	<u>0.06</u>	<u>0.62</u>	<u>0.05</u>	<u>0.56</u>	<u>0.03</u>



Comparison between study designs under diverse hypothesis (3)

Estimates assuming SD=pooled SD when SDA>SDP

10%		20%		30%		40%		50%	
N	Nr	N	Nr	N	Nr	N	Nr	N	Nr
190	215	209	238	228	264	248	287	269	313
CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)
<u>0.8</u>	<u>0.15</u>	<u>0.78</u>	<u>0.14</u>	<u>0.77</u>	<u>0.13</u>	<u>0.74</u>	<u>0.13</u>	<u>0.72</u>	<u>0.11</u>



Comparison between study designs under diverse hypothesis (4)

Estimates assuming $SDA > SDP$ when $SDA > SDP$									
10%		20%		30%		40%		50%	
N	Nr	N	Nr	N	Nr	N	Nr	N	Nr
214	232	252	278	294	321	339	370	388	421
CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)	CP	P(RI)
<u>0.85</u>	<u>0.19</u>	<u>0.84</u>	<u>0.19</u>	<u>0.85</u>	<u>0.21</u>	<u>0.85</u>	<u>0.22</u>	<u>0.85</u>	<u>0.21</u>

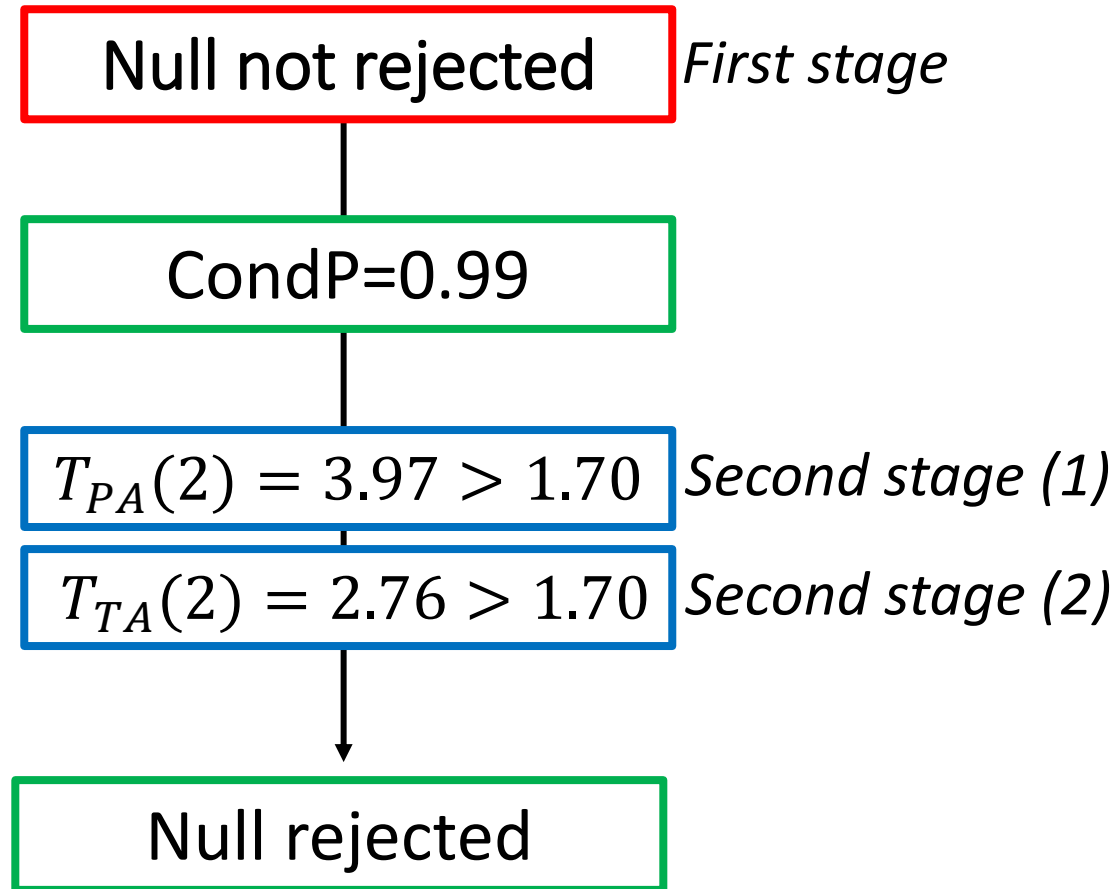


The real-data example (1)

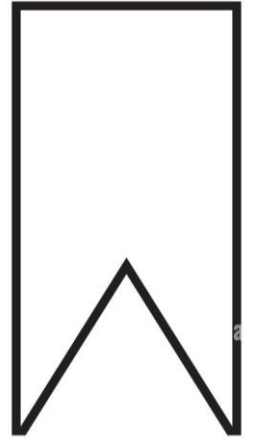
Labels	original	proposed	Difference (proposed - original)
Initial sample size	3276	2730	-546
Coverage Probability	0.876	0.811	-0.065
Average resampled sample size	3628	3191	-437
Probability to reject H0 at interim	0.112	0.062	-0.05
Probability to fall in unfavorable zone	0.164	0.206	0.042
Probability to fall in promising zone	0.209	0.231	0.022
Probability to fall in favorable zone	0.627	0.563	-0.064
WASS	2897	2471	-427



The real-data example (2)



Conclusions



- Take home messages
- Tutorial for the r-shiny web application



Take home message (1)



To our knowledge, no one has developed a parametrical multiple testing procedure to estimate the sample size for a three-arm noninferiority study with interim analysis for skewed/gamma distributed outcomes



We have used a **simulation approach** to empirically understand the **properties** of the statistical test under **deviation from normality**

Take home message (2)

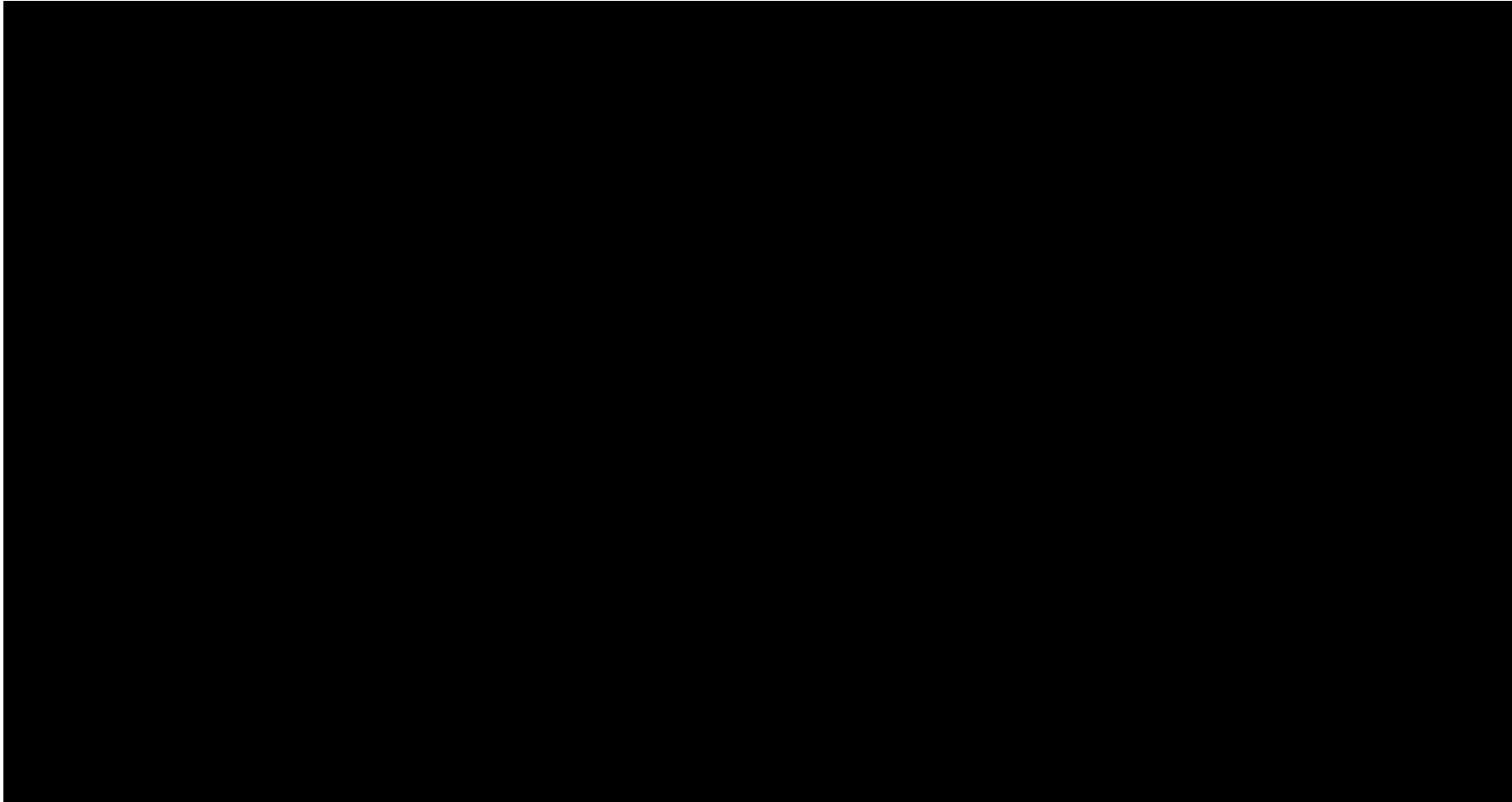


The first result was that **the original method, even under the assumption of non-normality, does not lose statistical power, assuming the variance of the three groups is equal or very similar**



We found that if **the discrepancy between the variances of the groups is considered**, using the proposed algorithm, we can estimate an initial sample size to achieve the desired power, **avoiding the risk of overestimation (saving patients) or underestimation (saving power)**, even in case of deviation from normality

Tutorial for the r-shiny web application



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Thank you very much for the
attention !

Do you have any question?



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The base-point of this study: the Lanyu Lei sample size algorithm

A group sequential design based on the promising zone of conditional power (2)

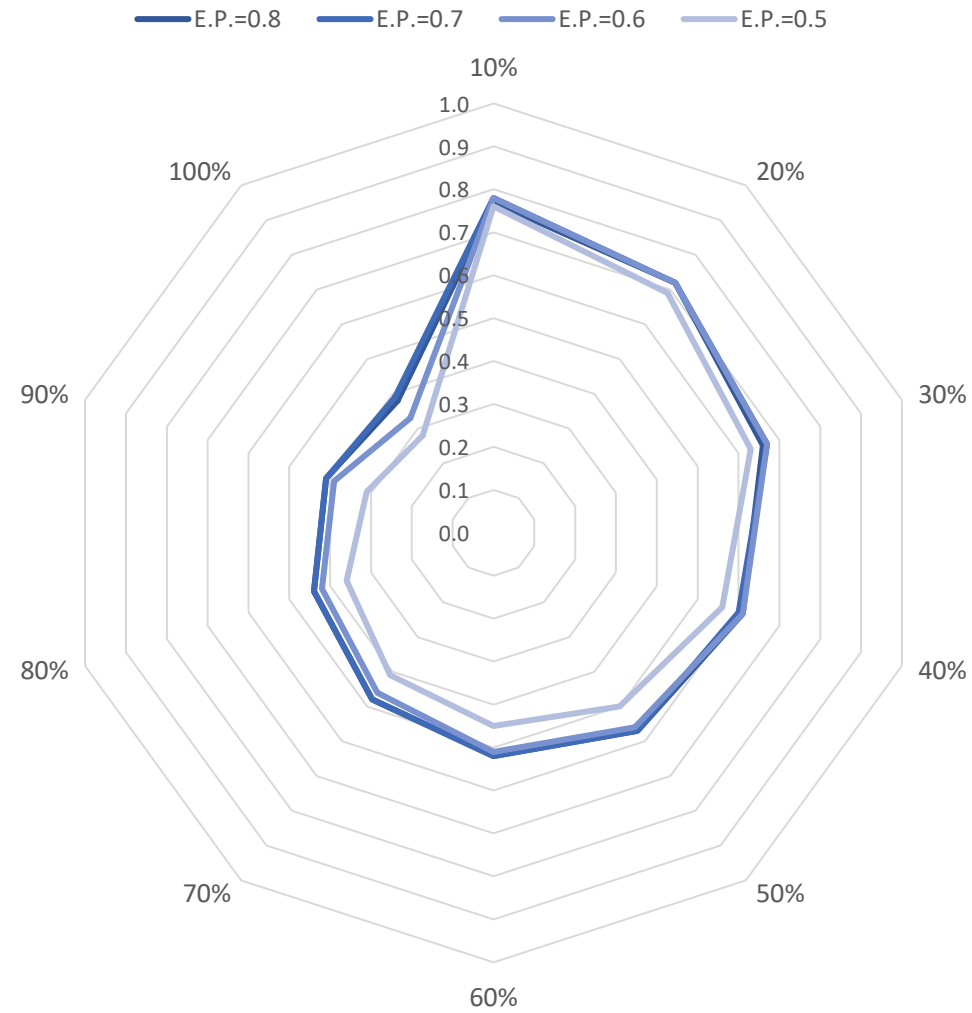
Table 4. The threshold of promising zones for the three-arm non-inferiority test in the two-stage SSR designs

n_{\max} (folds of original n)	Fraction of sample size at interim look ($n^{(1)}/n^{(2)}$)	Lower bound of promising zone
2	0.5	36%
2	0.75	33%
4	0.5	31%
4	0.75	30%
∞	0.5	31%
∞	0.75	30%

Note: The lower bounds were calculated for nominal initial power of 80%.



Comparison between study designs under diverse hypothesis (5)



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