

EqualStrength power analysis

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1 Introduction

1.1 Limitations

This analysis estimates the statistical power for EqualStrength tests across different sample sizes. Power is the probability of rejecting the null hypothesis when it is, in fact, false. In other words, the probability of correctly rejecting the hypothesis of non-discrimination when there is discrimination.

ignores the probability of the estimate being in the wrong direction or exaggerated [2]. 2 Main parameters

As with any power analysis, the scope of this analysis is limited to the statistical significance

of the test only. Apart from the general limitations of the NHST approach[1], this analysis

We estimate power as a function of a given i) sample size, ii) significance level, and iii) effect size.

▶ Code

For the sample size, we use a sequence of different sizes varying from 100 to 6000. The

significance level is determined by convention [3]: • Alpha: 0.05.

The effect size can be specified either from previous studies or by assuming a "minimum

effect that would be substantively important" [2] or interesting [4]. An overview of recent

studies indicate a substantial variation of effect sizes depending on the setting and country¹. Several studies also highlight that any postulation of effect sizes based on previous studies is severely limited by publication bias, population effect size heterogeneity, and model error [7], which can lead to "overly optimistic" expectations [4]. Here we posit the following callback rates for the main treatment: ▶ Code

• Majority group call-back rate: 0.4

▶ Code

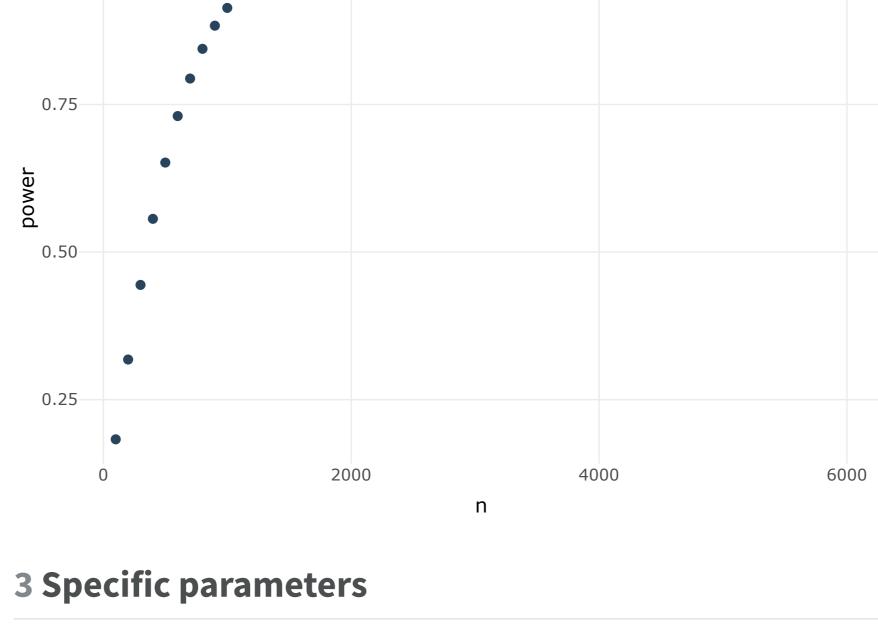
- Minority group call-back rate: 0.3
- We then use these rates to estimate the effect size (expressed as Cohen's h[3]):

• Effect size of the main treatment: -0.21.

2.1 Simple estimate (PWR) Based on these parameters, it is possible to estimate power assuming a basic design with two

groups of the same size. The package pwr [8] contains functions based on the formulas developed by Jacob Cohen[3] to calculate power. The following estimates are based on a power calculation for two proportions (with same sample sizes).

▶ Code 1.00



► Code

term between the two treatments.

The effect size of the second treatment is postulated to be the same as the one of the main treatment (-0.21). The interaction effect size is assigned as 25% or 50% the effect size of the main treatment, which results in the following values: • Interaction effect size (lower): 0.05.

EqualStrength research design. First we incorporate a second treatment and an interaction

Here we use three different approaches to incorporate specific features from the

3.1 Gelman, Hill, and Vehtari (GHV) In GHV[9], the authors highlight that it is necessary "4 times the sample size to estimate an

interaction that is the same size as the main effect" and estimate a 16 times larger sample for an interaction half the size of the main effect. Using the previous estimates from Section 2.1

• Interaction effect size (higher): 0.11.

and a power >80%, we would get a sample size of 3,200 for an interaction effect as large as the main effect and about 12,800 for an effect half the size.

3.2 Package InteractionPoweR (IPR) Using the package InteractionPoweR (IPR)[10] we can use the interaction effect size as a parameter for simulations.² It is assumed that the treatment variables are not correlated but this can be changed in the r.x1.x2 argument of the function. Code

Power of the interaction term

Power of the interaction term

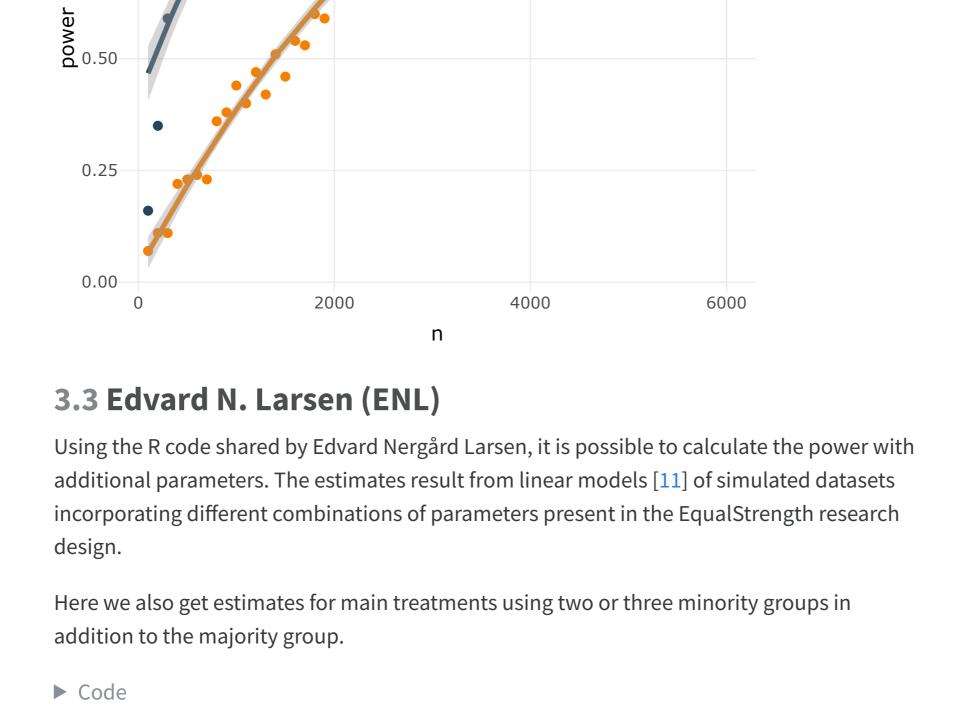
3 groups (2 minorities)

1.00

0.25

Interaction_ES 0.05 1.00 • 0.11





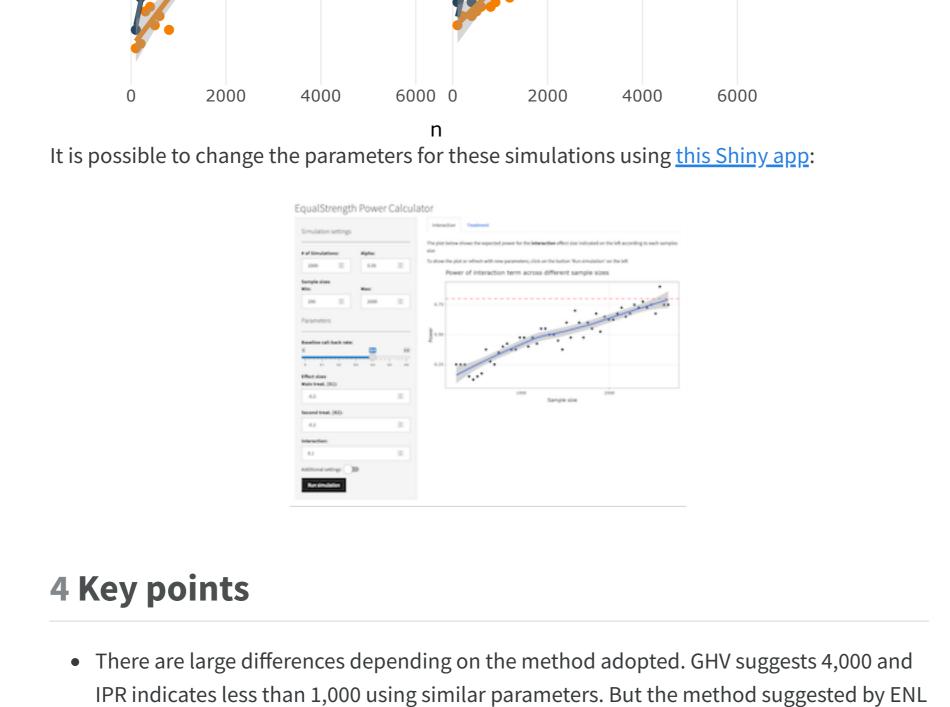
0.75 power

4 groups (3 minorities)

Interaction_ES

0.05

• 0.11



is the most appropriate as it incorporates attributes of the EqualStrength research

• Following this method, we estimate that at least 1,700 observations equally distributed

across 3 ethnic groups are necessary to get >= 80% power with a main effect size of -0.21

• After trying different parameters in the shiny app, the most influential component for the

0.075

0.15

Interaction

Min. sample 80%

600

600

3,100

sample size seems to be the effect size of the interaction term:

Main treat.

-0.3

-0.3

-0.3

Available from: https://doi.org/10.1177/1745691614551642

[Internet]. Elsevier; 1977 [cited 2023 Jul 4]. p. 1–17. Available from:

https://linkinghub.elsevier.com/retrieve/pii/B9780121790608500062

0.3 0.3

0.5

2.

3.

4.

6.

7.

Maj. call-back

design.

and an interaction effect size of 0.11.

0.3 -0.1 0.025 17,600 0.3 -0.1 0.05 5,600 -0.3 0.4 0.075 2,600 -0.3 0.4 1,100 0.15 0.4 -0.1 0.05 10,100

0.075

0.5 -0.3 0.15 11,00 0.5 -0.1 0.05 12,600 **5 References** McShane BB, Gal D, Gelman A, Robert C, Tackett JL. Abandon Statistical Significance. The 1. American Statistician [Internet]. 2019 Mar [cited 2023 Jul 4];73(sup1):235–45. Available from: https://doi.org/10.1080/00031305.2018.1527253

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doi=10.1037/xge0000920

- **Footnotes** 1. Looking at <u>some previous studies</u>, Ely found baseline callback rates ranging from 60-20%, CB rate
 - -17 to -6, and gender from +7 to +4. ←
- Method Adjusting Sample Effect Sizes for Publication Bias and Uncertainty. Psychological Science [Internet]. 2017 Nov [cited 2023 Jul 5];28(11):1547–62. Available from: for power analysis. 2017; Gelman A, Hill J, Vehtari A. Regression and Other Stories [Internet]. 1st ed. Cambridge University 9. Press; 2020 [cited 2023 Jul 5]. Available from:

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- differences from -31% (-0.31) to +7.3 (0.073). Ethnicity differences in some GEMM papers [5,6] runs from
- 2. The developers indicate that they would like to include 3-way interactions in a future version, but emphasise that "There are several steps in the analysis where we do not know exactly what to do for 3way interactions".←