

# Constructing Optimal Screening Designs for Effective Experimentation using Metaheuristics

Alan R. Vazquez

University of California, Los Angeles

[alanrvazquez@stat.ucla.edu](mailto:alanrvazquez@stat.ucla.edu)

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# Outline

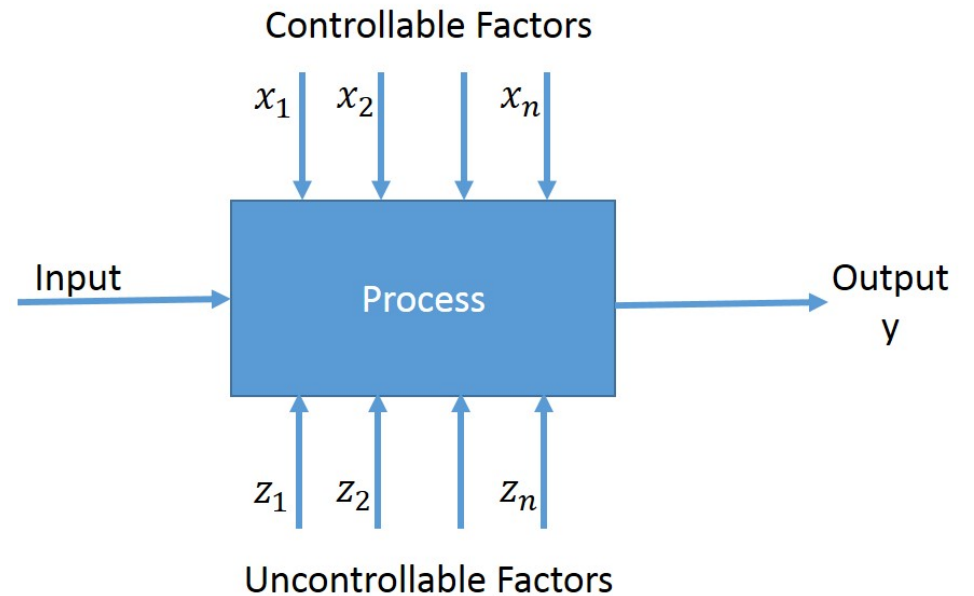
1. Introduction
2.  $Q_B$  criterion for evaluating experimental designs
3. A metaheuristic algorithm for constructing efficient designs
4. Results and Conclusions

# Experimental design

Structured plan for performing a series of tests of a system, a process or a product.

- Model:  $y = f(x_1, \dots, x_p) + \varepsilon$ .
- Interest is in finding economical experimental designs.

$f(x_1, \dots, x_p)$  includes main effects, two-factor interactions, quadratic effects, cubic effects, etc.



# Tuberculosis inhibition experiment

- Silva et al. (2016) conducted a study to develop a treatment that maximizes the percentage of inhibition of tuberculosis in infected human cells.
- The first stage of the study involved a screening experiment.
- 14 factors (drugs) at two levels (presence or absence).

## Goal:

Identify the influential main effects and two-factor interactions of the factors.

# Tuberculosis inhibition experiment

The first option for the experiment might be to test all the level combinations of the factors:  $2^{14} = 16,384$  tests!

Prior information:

1. Most of the 14 factors would be active.
2. Considerable number of active two-factor interactions.
3. Negligible three-factor and higher-order interactions.

How to select a fraction of the 16,384 level combinations?

For the experiment, the researchers chose an attractive experimental design with 14 factors and 128 test combinations (runs).

This design allowed the estimation of all main effects and all two-factor interactions with full precision.

In the end, however, only 8 main effects and 6 two-factor interactions were active.

### Research question:

Could we have identified the active effects with a smaller design?

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# Evaluating two-level designs

Example: Compare two-level designs with 24 runs and 7 factors.

**Design 1.**  
Folded-over  
Plackett-Burman  
design  
(Miller & Sitter, 2001)

X1	X2	X3	X4	X5	X6	X7
1	-1	1	1	1	-1	-1
1	-1	-1	-1	1	-1	1
1	-1	1	1	-1	1	1
1	1	1	-1	-1	-1	1
-1	1	-1	1	1	-1	1
-1	1	1	-1	1	1	1
-1	1	1	1	-1	-1	-1
-1	-1	-1	-1	-1	-1	-1
-1	-1	1	-1	1	1	-1
1	1	-1	-1	-1	1	-1
1	1	-1	1	1	1	-1
-1	-1	-1	1	-1	1	1
-1	1	-1	-1	-1	1	1
-1	1	1	1	-1	1	-1
-1	1	-1	-1	1	-1	-1
-1	-1	-1	1	1	1	-1
1	-1	1	-1	-1	1	-1
1	-1	-1	1	-1	-1	-1
1	-1	-1	-1	1	1	1
1	1	1	1	1	1	1
1	1	-1	1	-1	-1	1
-1	-1	1	1	1	-1	1
-1	-1	1	-1	-1	-1	1
1	1	1	-1	1	-1	-1

**Design 2.**  
Obtained from  
<http://neilsloane.com/hadamard/>

X1	X2	X3	X4	X5	X6	X7
-1	-1	1	1	1	1	1
1	-1	1	-1	-1	-1	-1
-1	1	1	1	1	1	-1
-1	-1	-1	-1	-1	-1	-1
1	1	1	1	-1	1	-1
1	1	-1	1	-1	1	1
1	1	1	1	1	-1	1
1	-1	-1	-1	-1	1	1
-1	1	-1	1	-1	-1	-1
1	-1	-1	1	-1	1	-1
-1	1	-1	-1	-1	-1	1
-1	-1	-1	-1	1	1	1
-1	1	1	-1	-1	1	-1
-1	-1	1	-1	1	-1	-1
1	-1	1	-1	1	1	-1
-1	-1	-1	1	1	1	1
-1	-1	1	1	-1	-1	1
1	1	-1	-1	1	-1	1
1	1	1	-1	1	-1	1
1	-1	1	1	-1	-1	1
-1	1	1	-1	-1	1	1
1	1	-1	-1	1	1	-1
-1	1	-1	1	1	-1	-1



# The $Q_B$ criterion

Measures the efficiency to estimate many potential models.

1. Maximal model including the intercept, all main effects and all two-factor interactions.
2. Sub-models of interest satisfy functional marginality.

$$y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \beta_{12} X_1 X_2 + \beta_{13} X_1 X_3 + \beta_{23} X_2 X_3 + \varepsilon$$

# The $Q_B$ criterion

Measures the efficiency to estimate many potential models.

1. Maximal model including the intercept, all main effects and all two-factor interactions.
2. Sub-models of interest satisfy functional marginality.
3.  $A_s$  criterion to measure the estimation efficiency of sub-models.

$$A_s = \sum_{i=1}^p \text{Var} \left( \hat{\beta}_i \right)$$

# The $Q_B$ criterion

Measures the efficiency to estimate many potential models.

## 4. Prior probabilities:

- $\pi_1$ : Active main effect.
- $\pi_2$ : Active interaction given that both of the main effects of the factors involved are active too.
- $\pi_3$ : Active interaction given that one of the main effects of the factors involved is active.

Under this framework, we can calculate the prior probability that sub-model is the best.

Li et al. (2006):  $\pi_1 = 0.41$ ,  $\pi_2 = 0.33$  and  $\pi_3 = 0.045$

# The $Q_B$ criterion

Weighted average of the  $A_s$  criterion over all sub-models of interest.

**Weights:** prior probability of a sub-model being the best.

Example I (cont.): 24-run 7-factor designs.

Consider  $\pi_1 = 0.41$ ,  $\pi_2 = 0.33$  and  $\pi_3 = 0.045$ .

**Design 1**

$$Q_B = 0.246$$

**Design 2**

$$Q_B = 0.255$$

Minimizing  $Q_B$  is equivalent to maximizing the estimation efficiency for the sub-models of the maximal model.

# An alternative expression

Let  $D$  be an  $n$ -run  $m$ -factor two-level design and  $T = DD^T$  with elements  $T_{ij}$ . We define  $M_k = \frac{1}{n^2} \sum_{i=1}^n \sum_{j=1}^n T_{ij}^k$  (Butler, 2003).

**Theorem:** For a maximal model including the intercept, all main effects and all two-factor interactions, minimizing the  $Q_B$  criterion is equivalent to minimizing

$$w_1 M_1 + w_2 M_2 + w_3 M_3 + w_4 M_4, \quad (1)$$

where  $w_k$  depends on  $\pi_1$ ,  $\pi_2$  and  $\pi_3$  only.

Computing the  $Q_B$  criterion using (1) is computationally cheap!

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# Metaheuristics

A metaheuristic is a high-level problem independent algorithmic framework that provides a set of guidelines or strategies to develop effective optimization algorithms (Sörensen & Glover, 2013).

## Local Search

Make small modifications to an existing single solution.

Variable Neighborhood Search  
Iterated Local Search

## Constructive

Build a single solution from scratch.

Greedy Randomize Adaptive Search  
Pilot method

## Population-Based

Work with a set of solutions which can communicate or be combined.

Genetic Algorithms  
Particle Swarm Optimization

# Local Search

- Local search algorithms perform local changes to an existing solution to attempt to find a better one.
- These algorithms, however, may get stuck in a locally optimal solution instead of the global optimum, since they do not examine all possible changes to the existing solution.

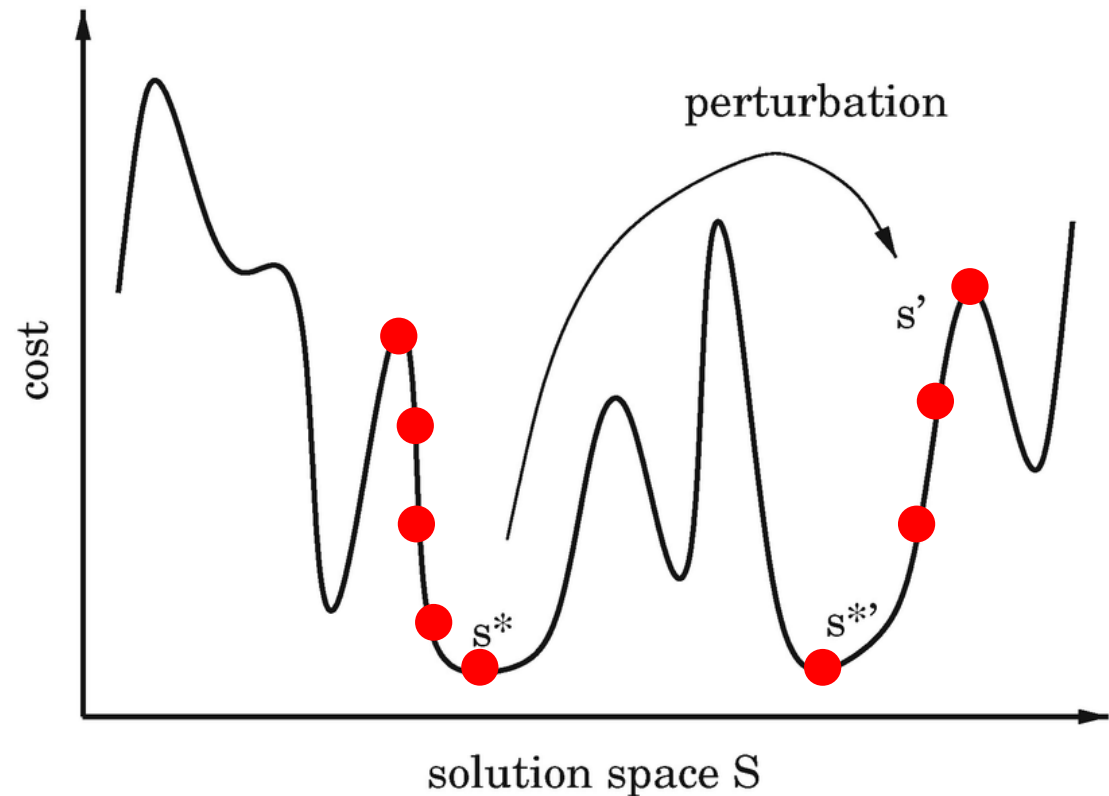
A local search metaheuristic uses an effective strategy to “escape” from this local optimum.



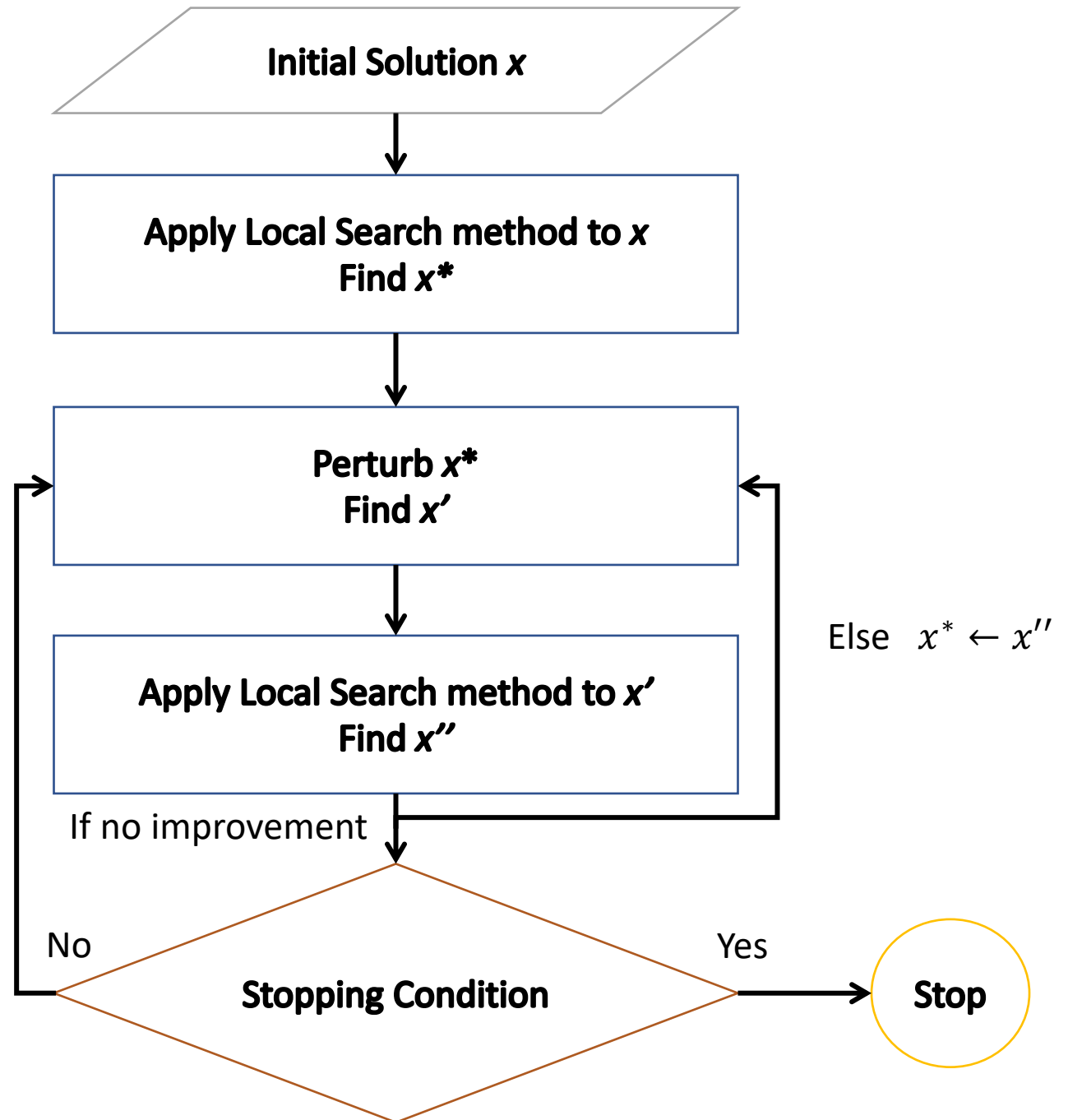
# Iterated Local Search (ILS)

## Principles:

- The solution obtained from a local search method is already quite good.
- Perform a small perturbation (modification) to that solution and start from there.



# Basic ILS



# Perturbation-Based Coordinate Exchange Algorithm

Constructs two-level designs that minimize the  $Q_B$  criterion.

## Building Blocks:

1. Local Search: Coordinate-exchange algorithm (Meyer & Nachtsheim, 1995).
2. Perturbation operator.

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.071$$

12 runs and 6 factors

$D =$

1	-1	1	1	1	-1
1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.064$$

$D =$

-1	-1	1	1	1	-1
1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.053$$

$D =$

-1	-1	1	1	1	-1
-1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.051$$

$D =$

-1	-1	1	1	1	-1
-1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
-1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.051$$

$D =$

-1	-1	1	1	1	-1
-1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
-1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1



# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.051$$

$D =$

-1	-1	1	1	1	-1
-1	-1	-1	1	-1	1
1	-1	1	1	-1	1
1	-1	-1	1	1	-1
1	1	1	-1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	-1	1
-1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	1	1	-1	1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.023$$

$D =$

-1	-1	-1	1	-1	-1
-1	-1	1	1	1	-1
1	-1	1	1	-1	1
1	1	-1	1	1	-1
1	-1	1	1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	1	1
-1	-1	-1	-1	1	1
-1	-1	-1	-1	-1	-1
1	-1	1	-1	-1	-1
1	1	-1	-1	-1	1

# Coordinate-Exchange Algorithm

- Local Search.
- One move: Sign switch a coordinate in the design  $D$ .

Example:

$$\pi_1 = 0.41, \pi_2 = 0.11, \pi_3 = 0.0$$

$$Q_B = 0.023$$

$D^* =$

-1	-1	-1	1	-1	-1
-1	-1	1	1	1	-1
1	-1	1	1	-1	1
1	1	-1	1	1	-1
1	-1	1	1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	1	1
-1	-1	-1	-1	1	1
-1	-1	-1	-1	-1	-1
1	-1	1	-1	-1	-1
1	1	-1	-1	-1	1

# Perturbation Operator

- Select coordinates at random and sign switch their elements.
- Tuning parameter  $\alpha$  denoted the percentage of coordinates that are affected.

$$D^* =$$

-1	-1	-1	1	-1	-1
-1	-1	1	1	1	-1
1	-1	1	1	-1	1
1	1	-1	1	1	-1
1	-1	1	1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
1	1	-1	1	1	1
-1	-1	-1	-1	1	1
-1	-1	-1	-1	-1	-1
1	-1	1	-1	-1	-1
1	1	-1	-1	-1	1

# Perturbation Operator

- Select coordinates at random and sign switch their elements.
- Tuning parameter  $\alpha$  denoted the percentage of coordinates that are affected.

Example:

$$\alpha = 0.05$$

$$D' =$$

-1	-1	-1	1	-1	-1
-1	-1	1	1	1	-1
1	-1	1	1	-1	1
1	1	-1	1	1	-1
1	-1	1	1	1	1
-1	1	1	-1	1	-1
-1	1	1	-1	-1	1
-1	1	-1	-1	1	1
-1	-1	-1	-1	1	1
-1	1	-1	-1	-1	1
1	-1	1	-1	-1	-1
1	1	-1	-1	-1	1

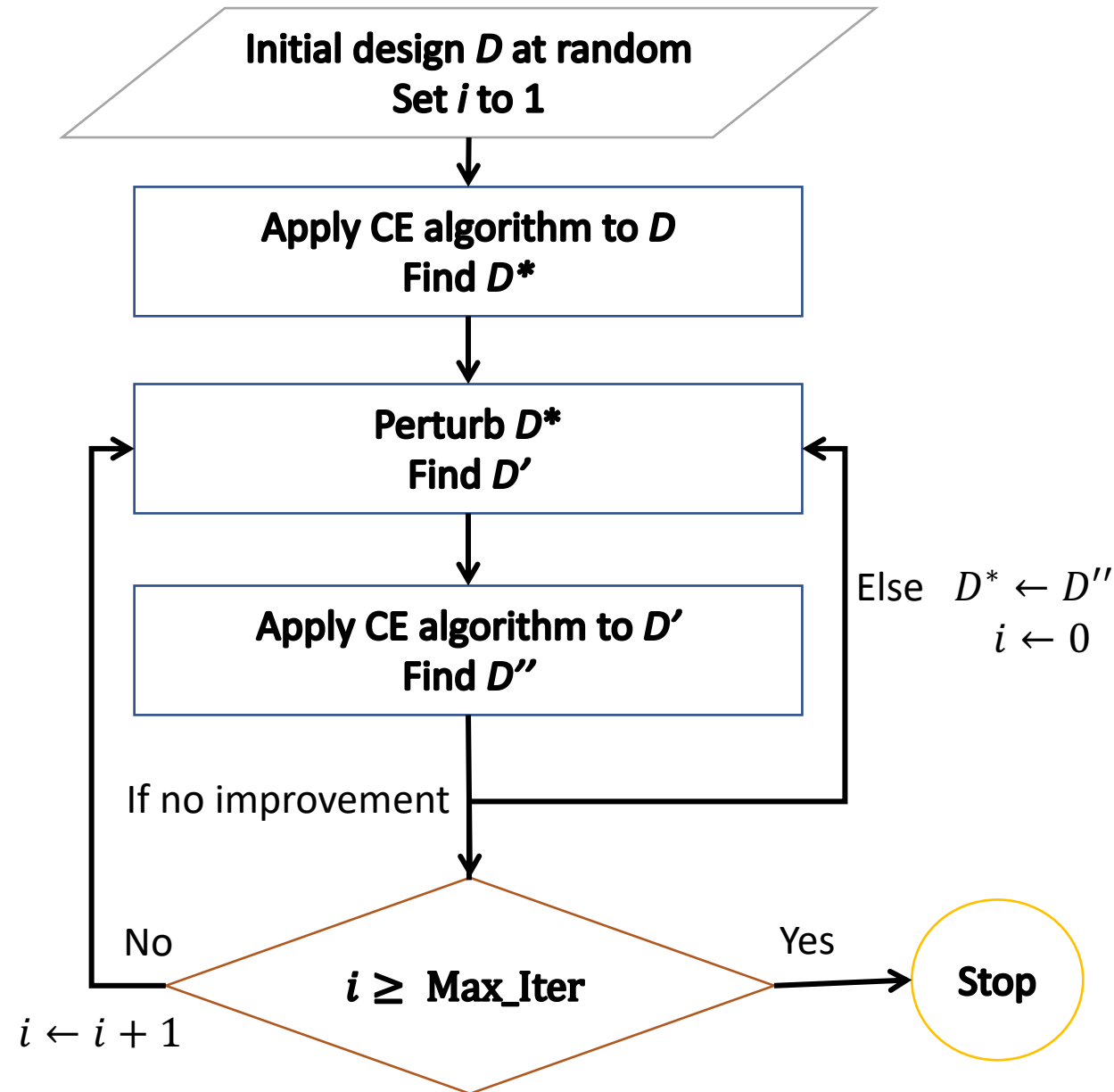
# PBCE algorithm in full

## Parameters:

**Max\_Iter**: maximum number of iterations without improvement.

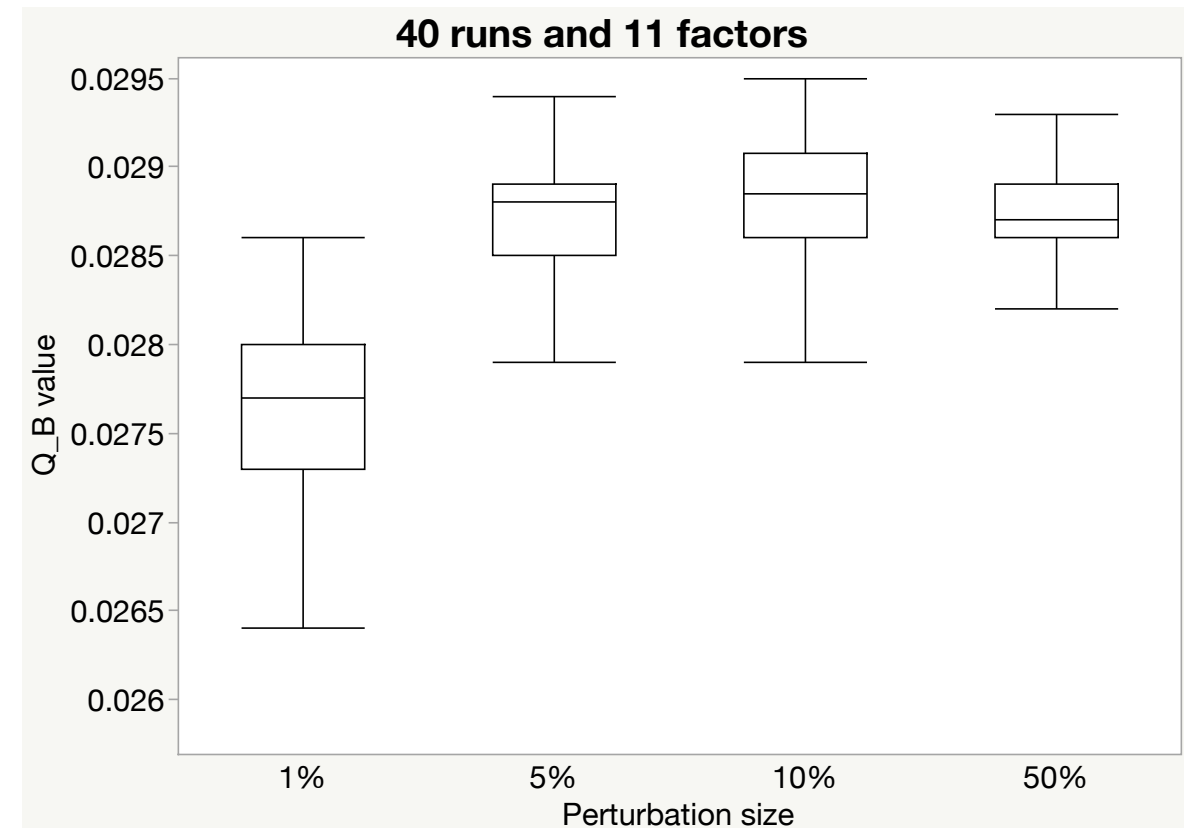
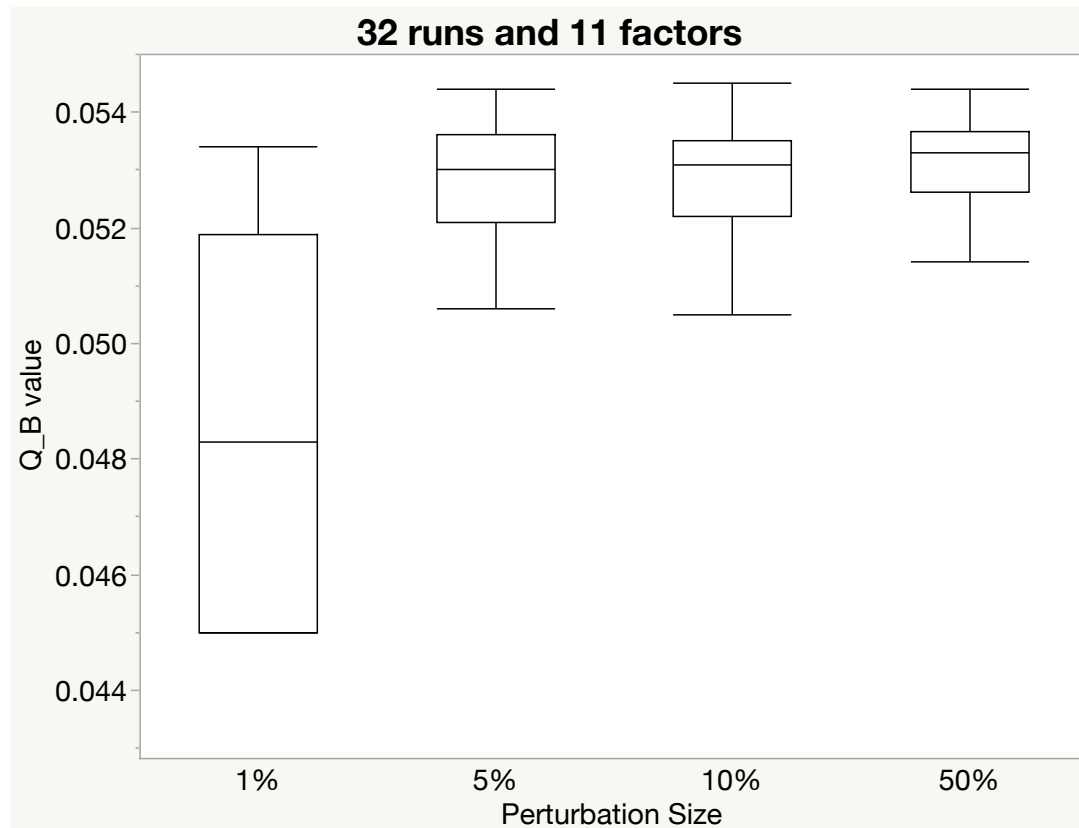
$\alpha$ : perturbation size

**Repetitions**: number of repetitions of the whole algorithm.



# Evaluation Perturbation Size ( $\alpha$ )

100  $Q_B$  values obtained using the PBCE algorithm with Max\_Iter = 100.



$$\pi_1 = 0.5, \pi_2 = 0.4, \pi_3 = 0.0$$

# Computing Times

10 repetitions of the PBCE algorithm with  $\alpha = 0.01$  and Max\_iter = 100.

Factors	Runs	Mean $\pm$ s.d. (seconds)
11	32	10.1 $\pm$ 2.7
11	40	18.8 $\pm$ 6.3
11	48	27.4 $\pm$ 11.0
14	40	11.18 $\pm$ 2.7
14	56	65.5 $\pm$ 22.4
14	64	101.7 $\pm$ 58.1

Implemented in Python v3.7.

Standard CPU with an Intel(R) Core(TM) i7 processor with 2.6Ghz and 16 GB of RAM.



# Benchmark Instances

We obtained benchmark instances involving 7 and 11 factors from Mee et al. (2017). We develop larger instances involving 16 factors.

Alternative algorithms:

- Coordinate-exchange algorithm with 1000 iterations (Meyer & Nachtsheim, 1995).
- Restricted columnwise-pairwise algorithm with 1000 iterations (Li, W. 2006).
- Mixed Integer Quadratic Programming with Gurobi and a maximum search time of 20 min.

Our PBCE algorithm involves an  $\alpha = 0.01$ , Max\_Iter = 100, and 5 repetitions.

# Results I

Factors	Runs	PBCE Algorithm	Coordinate- Exchange Algorithm	Restricted Columnwise- Pairwise Algorithm	Mixed Integer Quadratic Programming
$\pi_1 = 0.5, \pi_2 = 0.8$ and $\pi_3 = 0.0$					
7	16	<b>0.105</b>	<b>0.105</b>	<b>0.105</b>	<b>0.105</b>
	20	<b>0.0652</b>	<b>0.0652</b>	<b>0.0652</b>	<b>0.0652</b>
	24	<b>0.0333</b>	<b>0.0333</b>	0.0351	<b>0.0333</b>
	28	<b>0.0203</b>	<b>0.0203</b>	<b>0.0203</b>	<b>0.0203</b>
	32	<b>0.0075</b>	<b>0.0075</b>	<b>0.0075</b>	<b>0.0075</b>
$\pi_1 = 0.5, \pi_2 = 0.4$ and $\pi_3 = 0.0$					
11	20	<b>0.1726</b>	<b>0.1726</b>	0.1879	0.2154
	24	<b>0.0917</b>	<b>0.0917</b>	0.1189	0.1408
	32	<b>0.0478</b>	0.0531	0.0560	0.0664
	40	<b>0.0268</b>	0.0286	0.0318	0.0371
	48	<b>0.0161</b>	0.0168	0.0187	0.0235

Smaller the better

# Results II

Factors	Runs	PBCE Algorithm	Coordinate- Exchange Algorithm	Restricted Columnwise- Pairwise Algorithm	Mixed Integer Quadratic Programming
$\pi_1 = 0.41, \pi_2 = 0.33$ and $\pi_3 = 0.045$					
16	32	<b>0.0808</b>	<b>0.0808</b>	0.1374	-
	40	<b>0.0548</b>	0.0607	0.0794	-
	48	<b>0.0369</b>	0.0417	0.0519	-
	56	<b>0.0266</b>	0.0323	0.0356	-
	64	<b>0.0213</b>	0.0230	0.0256	-

Smaller the better

# Discussion of PBCE algorithm

- PBCE algorithm is the first dedicated algorithm to construct large designs that minimize the  $Q_B$  criterion.
- For large numbers of factors, our PBCE algorithm outperforms alternative algorithmic strategies.
- For up to 7 factors, our mixed integer quadratic programming approach is attractive since it can provide certificates of optimality.

Vazquez, A. R., Wong, W.-K., Goos, P. (2021). Constructing two-level  $Q_B$ -optimal designs for screening experiments using mixed integer programming and heuristic algorithms. In preparation.

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# Alternative designs for the tuberculosis inhibition experiment

- We constructed alternative 14-factor designs with 64, 80, 96 and 112 runs.
- Using the PBCE algorithm, we generated designs which optimize the  $Q_B$  criterion with  $\pi_1 = 0.41$ ,  $\pi_2 = 0.33$  and  $\pi_3 = 0.045$  (Li et al., 2006).
- We evaluate the performance of these designs using a simulation study.

# Available designs in the literature

- 14-factor Bayesian D-optimal designs with 64, 80, 96 and 112 runs (DuMouchel & Jones, 1994).
- 14-factor D-optimal design with 112 runs (Atkinson et al., 2007).
- 14-run design with 128 runs (Silva et al. , 2016).

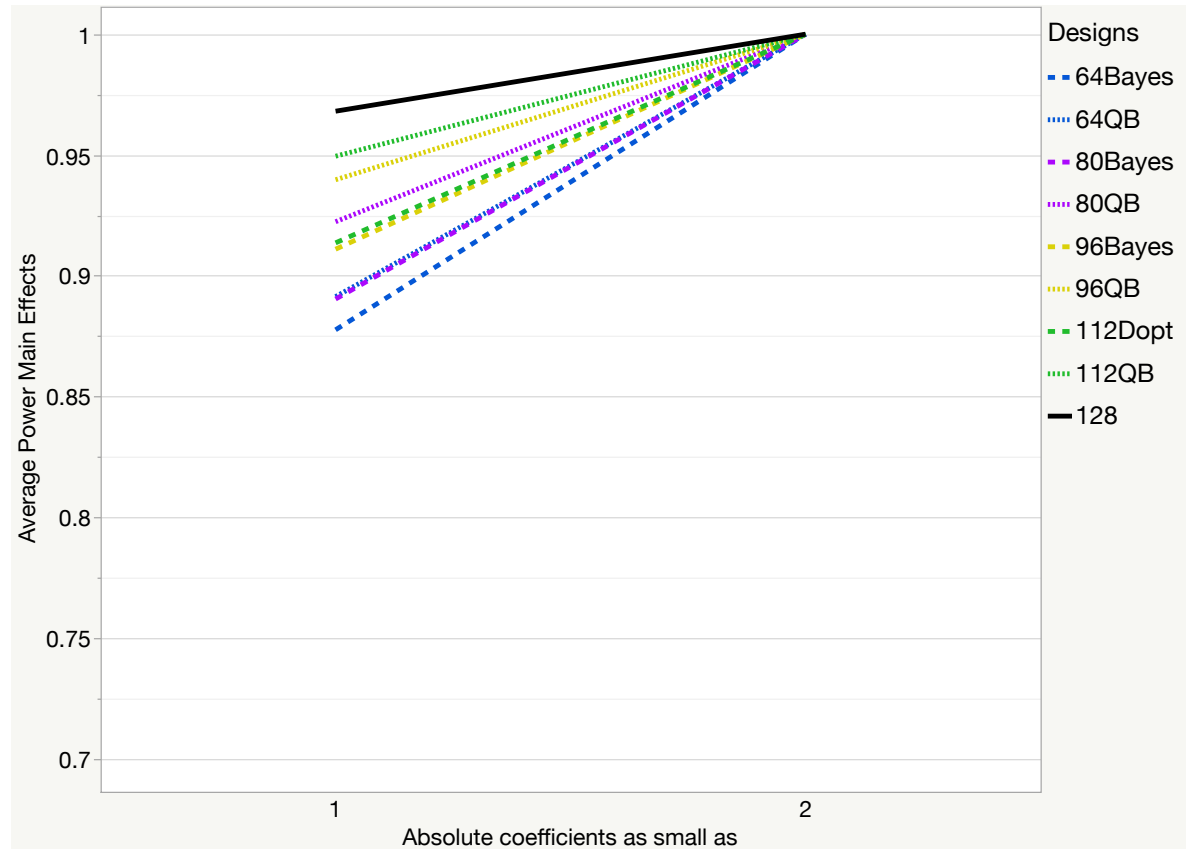
# Simulation Study

1. We randomly selected 8 main effects and 6 two-factor interactions (satisfying weak effect heredity) as active.
2. We obtained coefficients for the active effects by adding a 1 or a 2 to an exponentially distributed random number. A '+' or '-' was randomly assigned.
3. The coefficients for the inactive effects were drawn from  $N(0, 0.25^2)$ .
4. We simulated response vectors with residuals following  $N(0, 1)$ .
5. We used the Dantzig selector (Candes & Tao, 2007) to identify the active effects.

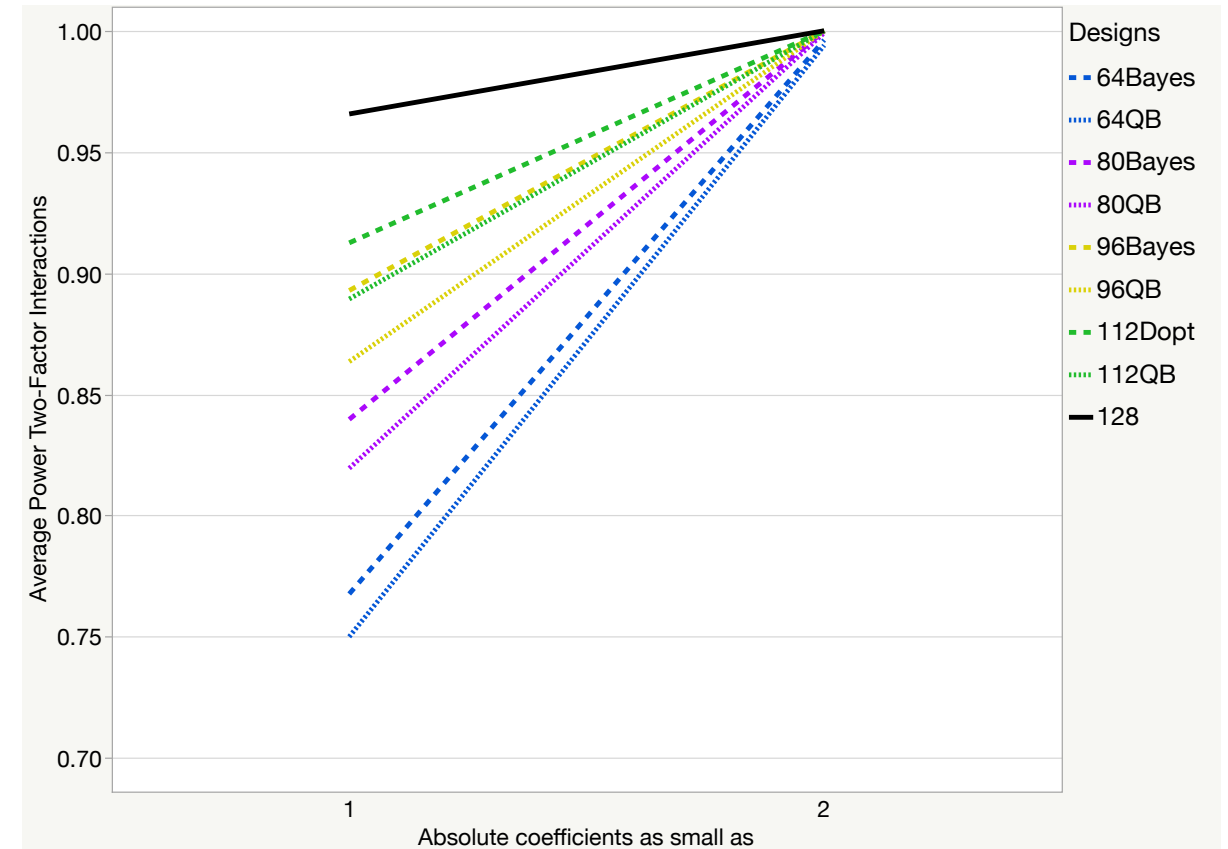
**1,000 simulations**



## Main Effects



## Two-Factor Interactions



Power: Proportion of active effects that are successfully detected.

# Conclusions

- The tuberculosis inhibition experiment would have benefited from a smaller design.
- Metaheuristics allow the development of effective algorithms to construct attractive experimental designs.
- Two-level designs can be used to find the right hyperparameter values of machine learning algorithms (Choueki et al., 1997; Lujan-Moreno, 2018).
- Future research: Explore other metaheuristics such as particle swarm optimization to construct  $Q_B$ -optimal designs with two or more levels.

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