

Design and Analysis of Computer Experiments

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November 2020

Outline

- 1 Introduction
- 2 Modeling Computer Experiments
- 3 Designs for Computer Experiments
- 4 Uniform Projection Designs
- 5 Construction Methods
- 6 References

What are computer experiments?

- Computer experiments are increasingly being used to explore the behavior of complex physical systems.
- A computer model is a **large** computer code that implements a **complex** mathematical model of a physical process.
e.g., simultaneous differential solver, finite element analysis computational fluid dynamics.

A typical engineering model (Page 1 of 3)

$$\begin{aligned}
 & \rho_b A_b \frac{\partial^2 w}{\partial t^2} + E_b A_b \frac{\partial^4 w}{\partial x^4} \\
 & + \left[(\rho_c A_c + \rho_e A_e) \frac{\partial^2 w}{\partial t^2} + \rho_c A_c \left(\frac{t_b + t_c}{2} \right) \left(\frac{\partial^3 u_b}{\partial x \partial t^2} - \frac{t_b + t_c}{2} \frac{\partial^4 w}{\partial x^2 \partial t^2} - \frac{t_c}{2} \frac{\partial^3 \beta}{\partial x \partial t^2} \right) \right. \\
 & + \rho_e A_e \left(\frac{\partial^3 u_b}{\partial x \partial t^2} - a \frac{\partial^4 w}{\partial x^2 \partial t^2} + t_c \frac{\partial^3 \beta}{\partial x \partial t^2} \right) + C_{11}^0 I_c \frac{\partial^4 w}{\partial x^4} - E_c A_c a \left(\frac{\partial^3 u_b}{\partial x^3} - a \frac{\partial^4 w}{\partial x^4} + t_c \frac{\partial^3 \beta}{\partial x^3} \right) \left. \right] [H(x - x_1) - H(x - x_2)] \quad (1) \\
 & + \left[\rho_c A_c \left(\frac{t_b + t_c}{2} \right) \left(\frac{\partial^3 u_b}{\partial t^3} - \frac{t_b + t_c}{2} \frac{\partial^3 w}{\partial x \partial t^2} + \frac{t_c}{2} \frac{\partial^3 \beta}{\partial t^3} \right) + \rho_e A_e a \left(\frac{\partial^3 u_b}{\partial t^3} - a \frac{\partial^3 w}{\partial x \partial t^2} + t_c \frac{\partial^3 \beta}{\partial t^3} \right) \right. \\
 & \left. + 2C_{11}^0 I_c \frac{\partial^3 w}{\partial x^3} - 2E_c A_c a \left(\frac{\partial^2 u_b}{\partial x^2} - a \frac{\partial^3 w}{\partial x^2} + t_c \frac{\partial^2 \beta}{\partial x^2} \right) \right] [\delta(x - x_1) - \delta(x - x_2)] \\
 & + [C_{11}^0 I_c \frac{\partial^2 w}{\partial x^2} - E_c A_c a \frac{\partial u_b}{\partial x} - a \frac{\partial^2 w}{\partial x^2} - t_c \frac{\partial \beta}{\partial x} + b d_{11} E_c V(t)] [\delta'(x - x_1) - \delta'(x - x_2)] = f(x, t)
 \end{aligned}$$

$$\begin{aligned}
 & \rho_b A_b \frac{\partial^2 u_b}{\partial t^2} - E_b A_b \frac{\partial^2 u_b}{\partial x^2} \\
 & + \left[\rho_c A_c \left(\frac{\partial^2 u_b}{\partial t^2} - \frac{t_b + t_c}{2} \frac{\partial^3 w}{\partial x \partial t^2} - \frac{t_c}{2} \frac{\partial^3 \beta}{\partial t^2} \right) + \rho_e A_e \left(\frac{\partial^2 u_b}{\partial t^2} - a \frac{\partial^3 w}{\partial x \partial t^2} + t_c \frac{\partial^2 \beta}{\partial t^2} \right) \right. \\
 & \left. - E_c A_c \left(\frac{\partial^2 u_b}{\partial x^2} - a \frac{\partial^3 w}{\partial x^2} + t_c \frac{\partial^2 \beta}{\partial x^2} \right) \right] [H(x - x_1) - H(x - x_2)] \\
 & + \left[-E_c A_c \left(\frac{\partial u_b}{\partial x} - a \frac{\partial^2 w}{\partial x^2} - t_c \frac{\partial \beta}{\partial x} \right) + b d_{11} E_c V(t) \right] [\delta(x - x_1) - \delta(x - x_2)] = 0
 \end{aligned} \quad (2)$$

$$\begin{aligned}
 & \left[\rho_c A_c \left(\frac{t_c}{2} \left(\frac{\partial^2 u_b}{\partial t^2} - \frac{t_b + t_c}{2} \frac{\partial^3 w}{\partial x \partial t^2} + \frac{t_c}{2} \frac{\partial^3 \beta}{\partial t^2} \right) + \rho_e A_e t_c \left(\frac{\partial^2 u_b}{\partial t^2} - a \frac{\partial^3 w}{\partial x \partial t^2} + t_c \frac{\partial^2 \beta}{\partial t^2} \right) \right. \right. \\
 & \left. \left. + A_c (G + \beta) - E_c A_c t_c \left(\frac{\partial^2 u_b}{\partial x^2} - a \frac{\partial^3 w}{\partial x^2} + t_c \frac{\partial^2 \beta}{\partial x^2} \right) \right] [H(x - x_1) - H(x - x_2)] \quad (3)
 \end{aligned}$$

Computer Experiments

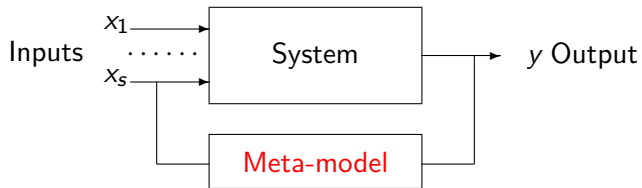


Figure 1: Computer experiment

Characteristics of computer experiments

- Mostly **deterministic** (**lack of random error**)
- May take hours or even days to produce a single output
- Many input variables
- The performance of the predictor depends upon the choice of the training data (**design**).

Principles in traditional DOE are irrelevant

- Replication
- Blocking
- Randomization

Modeling Computer Experiments: Kriging

For $x \in \mathcal{R}^m$, treat the deterministic response $y(x)$ as a realization of a stochastic process

$$Y(x) = \sum_{j=1}^k \beta_j f_j(x) + Z(x),$$

where $f_j(x)$ are known functions, β_j are unknown parameters and $Z(\cdot)$ is a **Gaussian process** with mean 0 and covariance

$$\text{cov}(Z(w), Z(x)) = \sigma^2 R(w, x).$$

- This is the Kriging model used in spatial statistics.
- Also called Gaussian process model in Machine Learning.

R packages: DiceKriging, kerngp, etc.

Prediction

Given a design $S = \{s_1, \dots, s_n\}$ and data $y_S = (y(s_1), \dots, y(s_n))^T$. Consider the linear predictor

$$\hat{y}(x) = c(x)^T y_S.$$

Frequentists replace y_S by the random vector $Y_S = \{Y(s_1), \dots, Y(s_n)\}^T$, and compute the MSE.

The **Best Linear Unbiased Predictor** (BLUP): choose $c(x)$ to minimize

$$\text{MSE} [\hat{y}(x)] = E[c(x)^T Y_S - Y(x)]^2$$

subject to

$$E[\hat{y}(x)] = E[c(x)^T Y_S] = E[Y(x)]$$

Kriging model: $Y(x) = f(x)^T \beta + Z(x)$, where

$$f(x) = (f_1(x), \dots, f_k(x))^T, \beta = (\beta_1, \dots, \beta_k)^T$$

In matrix form:

$$Y_S = F\beta + Z, \text{ cov}(Z) = \sigma^2 R$$

$$F = (f(s_1), \dots, f(s_n))^T = (f_j(s_i))_{n \times k}$$

$$R = (R(s_i, s_j))_{n \times n}$$

$$r(x) = (R(s_1, x), \dots, R(s_n, x))^T$$

The generalized LS estimate and BLUP are

$$\hat{\beta} = (F^T R^{-1} F)^{-1} F^T R^{-1} Y_S$$

$$\hat{y}(x) = f(x)^T \hat{\beta} + r(x)^T R^{-1} (Y_S - F \hat{\beta})$$

The GP interpolates the observed data: $\hat{y}(s_i) = y(s_i)$ for $s_i \in S$.

Correlation Functions

The correlation $R(w, x)$ has to be specified. Commonly used functions:

$$R(w, x) = \prod \exp(-\theta_j |w_j - x_j|^{p_j}), \quad 0 < p_j \leq 2,$$

$$R(w, x) = \prod K(|w_j - x_j|; \theta_j)$$

where $K()$ is Matérn correlation function with parameter $\nu = 5/2$.

$$K(h; \theta) = \left(1 + \frac{\sqrt{5}h}{\theta} + \frac{5h^2}{3\theta^2} \right) \exp \left(-\frac{\sqrt{5}h}{\theta} \right).$$

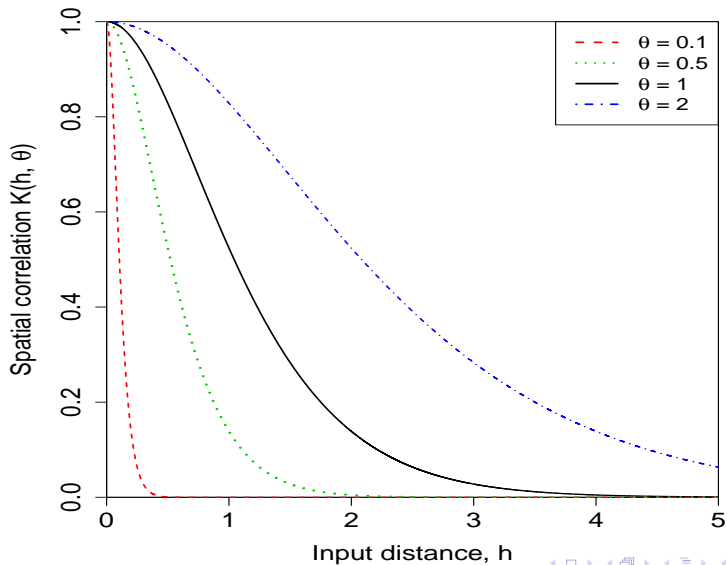
The correlation parameters (e.g., θ_j , p_j) need to be specified or estimated (by MLE or cross validation)

Given the correlation parameters, the MLEs are

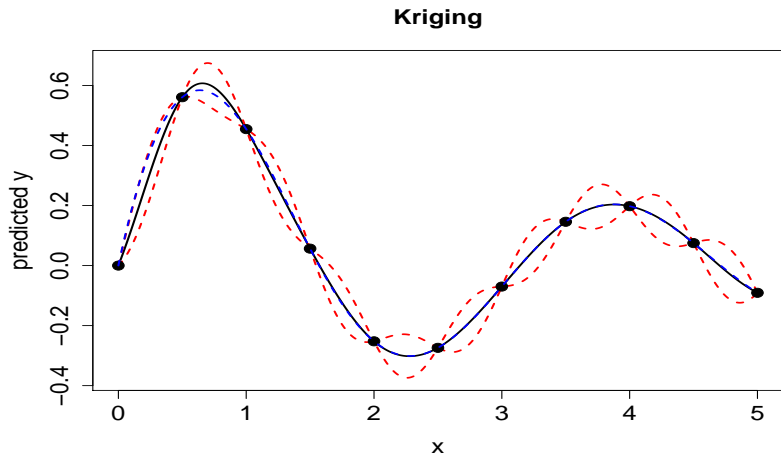
$\hat{\beta}$ = generalized l.s. estimate

$$\hat{\sigma}^2 = \frac{1}{n} (Y_S - F\hat{\beta})' R^{-1} (Y_S - F\hat{\beta})$$

Examples of Matérn $\nu = 5/2$ correlation functions

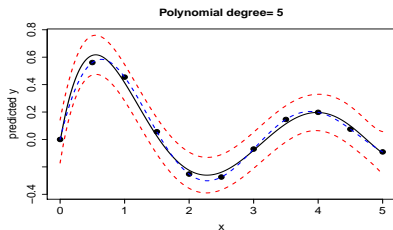
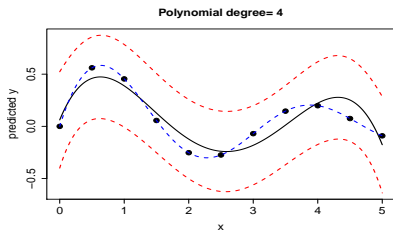
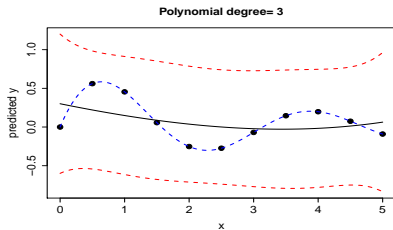
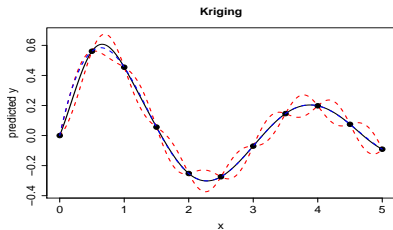


A toy example: Kriging



Data: $y = \sin(2x)/(1 + x)$; Kriging: $Y = \mu + Z(x)$.

A toy example: Kriging vs Polynomial models



Data: $y = \sin(2x)/(1+x)$; Kriging: $Y = \mu + Z(x)$.

Designs for Computer Experiments

- Constructing a “good” design is crucial for the success of a computer experiment.
- A “good” design should be **space-filling** (i.e., cover as much space as possible), and have **good projection properties**.
 - **Latin hypercube designs** (LHD) [McKay et al. (1979)]
 - **Maximin and minimax distance designs** [Johnson et al. (1990)]
 - **Orthogonal Array-based designs** [Owen (1992), Tang (1993), He and Tang (2013, 2014)]
 - **Uniform designs** [Fang et al. (2006)]
 - **Maximum projection designs** [Joseph et al. (2015)]
 - **Uniform projection designs** [Sun et al. (2019)]
- Optimality criteria: maximin distance, minimax distance, column-orthogonality, uniformity (discrepancy) etc.
- R packages: lhs, LHD, SLHD, UniDOE, MaxPro, etc.

Design Criteria

Let $\hat{y}(x)$ be BLUE of $y(x)$ given a design $S = \{s_1, \dots, s_n\}$.

- Integrated Mean Squared Error (IMSE)

$$\min_S : \int_{\mathcal{X}} \text{MSE} [\hat{y}(x)] \phi(x) dx,$$

where $\phi(x)$ is a given weight function.

- Maximum Mean Squared Error (MMSE)

$$\min_S : \max_{x \in \mathcal{X}} \text{MSE} [\hat{y}(x)].$$

- Entropy (Gaussian process)

$$\max_S : \det(R) = \det(R(s_i, s_j)).$$

- Maximin distance criterion:

$$\max_S : \min_{i < j} d(s_i, s_j)$$

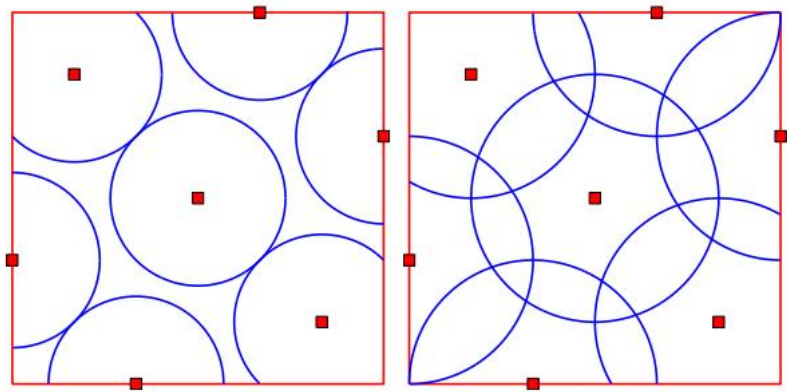


Figure 2: Maximin LHD (left) and Minimax LHD (right) with $n = 7$ and $m = 2$

Maximin distance designs

- For an (n, s^m) design $D = (x_{ik})_{n \times m}$,

$$d_p(x_i, x_j) = \sum_{k=1}^m |x_{ik} - x_{jk}|^p,$$

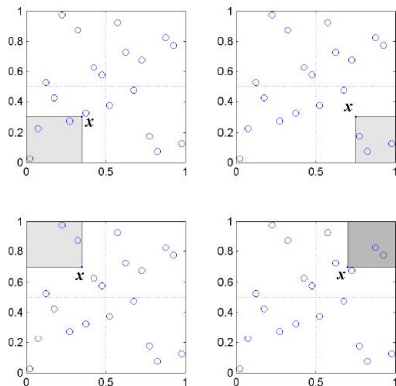
- Define the L_p -distance of D as

$$d_p(D) = \min\{d_p(x_i, x_j), 1 \leq i < j \leq n\}$$

- Maximin distance design: maximize $d_p(D)$ among all designs
- Most constructions are based on stochastic algorithms:
 - Morris and Mitchell (1995), Joseph and Hung (2008), Ba, Myers and Brennenman (2015, R package SLHD), etc.
 - Flexible but are not effective for large designs.
- Low-dimensional projections may not be space-filling.
 - Saturated $OA(n, 2^m)$'s are maximin distance designs when $m = n - 1$ (Xu 1999).

Uniform designs

Idea: choose design points from the design region with empirical distribution as “uniform” as possible (Fang et al, 2006).



Uniform Designs and Centered L_2 -Discrepancy

For an $n \times m$ design D over $[0, 1]^m$,

$$\text{Disc}(D) = \left\{ \int_{[0,1]^m} \left| \text{Vol}(J(a_x, x)) - \frac{N(D \cap J(a_x, x))}{n} \right|^2 dx \right\}^{1/2}.$$

The (squared) centered L_2 -discrepancy is defined by

$$\text{CD}(D) = \left\{ \sum_{u \subseteq \{1:m\}} |\text{Disc}(D_u)|^2 \right\},$$

where D_u is the projected design of D onto dimensions indexed by the elements of u .

- Uniform designs may have poor projections in lower dimensional spaces.

Uniform Projection Designs: A New Class of Space-Filling Designs

- Focus on 2-dim projection uniformity
- *Uniform projection criterion* (Sun, Wang and Xu, 2019, Annals of Statistics)

$$\phi(D) = \frac{2}{m(m-1)} \sum_{|u|=2} CD(D_u), \quad (1)$$

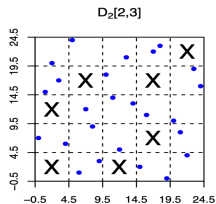
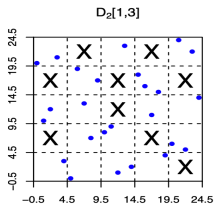
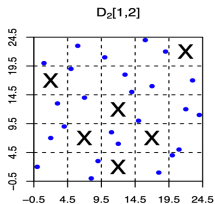
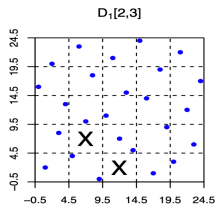
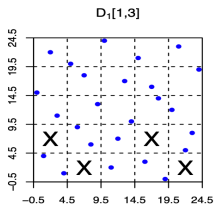
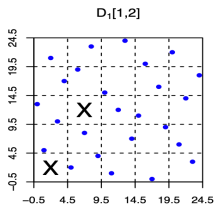
- A design achieving the minimum $\phi(D)$ value is a **uniform projection design** (UPD).
- The discrepancy has an analytical expression; for $D = (z_{ik})$ over $[0, 1]^m$:

$$\begin{aligned} CD(D) = & \frac{1}{n^2} \sum_{i=1}^n \sum_{j=1}^n \prod_{k=1}^m \left(1 + \frac{1}{2}|z_{ik}| + \frac{1}{2}|z_{jk}| - \frac{1}{2}|z_{ik} - z_{jk}| \right) \\ & - \frac{2}{n} \sum_{i=1}^n \prod_{k=1}^m \left(1 + \frac{1}{2}|z_{ik}| - \frac{1}{2}|z_{ik}|^2 \right) + \left(\frac{13}{12} \right)^m. \end{aligned}$$

Why we need a new criterion? Four 25×3 LHDs

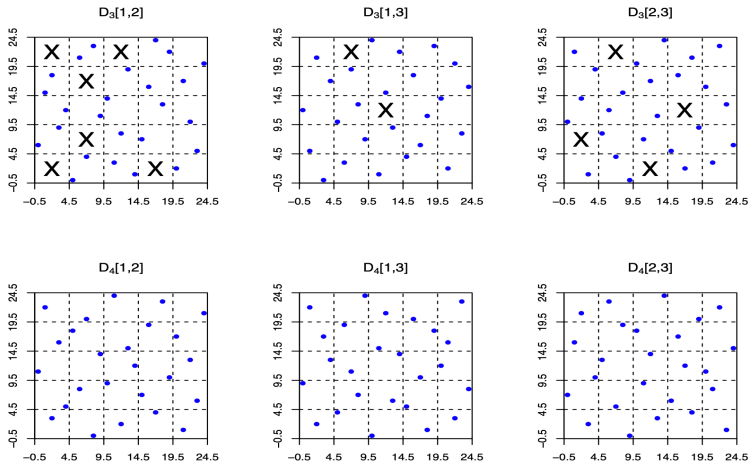
Uniform D_1			Maximin D_2			MaxPro D_3			UPD D_4		
18	16	14	0	2	20	0	6	12	2	3	2
19	9	0	1	20	10	1	15	5	4	5	13
11	1	2	2	7	12	2	18	21	0	11	9
16	20	3	3	13	21	3	9	0	3	16	17
20	22	12	4	9	3	4	12	17	1	22	22
14	7	10	5	19	0	5	0	10	8	0	7
4	17	1	6	23	19	6	21	3	6	8	19
12	12	7	7	14	13	7	4	19	9	14	24
10	15	24	8	0	7	8	23	13	5	18	4
22	14	5	9	3	17	9	11	7	7	20	11
2	21	22	10	21	8	10	14	24	12	2	21
15	11	21	11	8	9	11	3	1	10	9	0
1	5	4	12	6	1	12	8	15	14	12	14
3	10	11	13	18	23	13	19	9	13	15	6
23	3	8	14	15	2	14	1	22	11	24	15
0	13	15	15	10	18	15	7	4	17	4	10
8	23	6	16	24	16	16	16	18	15	7	5
7	8	18	17	16	11	17	24	6	19	10	18
9	4	13	18	1	15	18	13	11	16	19	20
6	19	9	19	22	4	19	22	23	18	23	1
24	18	19	20	4	6	20	2	14	21	1	16
21	6	23	21	5	24	21	17	2	23	6	23
13	24	17	22	12	5	22	10	20	22	13	3
17	0	16	23	17	22	23	5	8	20	17	12
5	2	20	24	11	14	24	20	16	24	21	8

Bivariate projections of Uniform D_1 and Maximin D_2



Note: 'X' means that there are no points in the grid.

Bivariate projections of MaxPro D_3 and UPD D_4



Note: 'X' means that there are no points in the grid.

Some Theoretical Results

Theorem 1

For a balanced (n, s^m) design D and any $2 \leq k \leq m$,

$$\frac{1}{\binom{m}{k}} \sum_{|u|=k} \phi(D_u) = \phi(D),$$

where D_u is the projected design onto k factors indexed by u .

- UPDs have good space-filling properties not only in two dimensions, but also in all dimensions.

Some Theoretical Results

Theorem 2

For a balanced (n, s^m) design $D = (x_{ik})$,

$$\phi(D) = \frac{g(D)}{4m(m-1)n^2s^2} + C(m, s), \quad (2)$$

where

$$g(D) = \sum_{i=1}^n \sum_{j=1}^n d_1^2(x_i, x_j) - \frac{2}{n} \sum_{i=1}^n \left(\sum_{j=1}^n d_1(x_i, x_j) \right)^2 \quad (3)$$

- $\phi(D)$ is a function of pairwise L_1 -distances of the rows of D .
- An equidistant design under the L_1 -distance is a UPD.

Application: Design and Modeling Comparison

- A 3-drug combination experiment on lung cancer (Al-Shyoukh et al. 2011; Xiao, Wang and Xu 2019).
- A 512-run and 8-level full factorial design to study 3 drugs.
- The response was the ATP level of the cells after the drug treatments.
- Kriging model with noise: $y(x) = \mu + Z(x) + \epsilon$

Table 1: Comparison of $1000 \times \text{MSE}$ for different models and designs

	Normal Cell				Cancer Cell			
	D ₅₁₂	RD ₈₀	MPD ₂₅	UPD ₂₅	D ₅₁₂	RD ₈₀	MPD ₂₅	UPD ₂₅
Kriging	0.002	0.21	0.62	0.22	0.003	0.37	1.87	0.21
NN	0.37	1.28	3.12	1.79	0.47	1.57	4.10	2.93
Polynomial	0.48	1.16	3.22	0.74	2.98	6.77	10.04	4.42

RD₈₀: Random 80-run design; MPD₂₅: MaxPro 25-run designs.

Comparison of projection properties

We compare four LHD(19, 18)'s:

- 1 The uniform design is from the uniform design website(UD)
- 2 The maximin distance design via R package SLHD (Ba, Myers and Brenneman, 2015, Technometrics).
- 3 The maximum projection (MaxPro) design were constructed via R package MaxPro (Joseph et al., 2015, Biometrika)
- 4 The uniform projection design (UPD): E_b .

We ran R commands `maximinSLHD` (with slice parameter $t = 1$) and `MaxProLHD` 100 times with default settings and chose the best designs.

Comparison of projection properties

Four criteria will be used in the comparison:

- 1 minimum Euclidean distance
- 2 maximum projection criterion (Joseph et al. 2015)

$$\psi(D) = \left\{ \frac{1}{\binom{n}{2}} \sum_{i=1}^{n-1} \sum_{j=i+1}^n \frac{1}{\prod_{k=1}^m (x_{ik} - x_{jk})^2} \right\}^{1/m}$$

- 3 relative maximum centered L_2 -discrepancy (CD)
- 4 maximum correlation ρ_{ave} .

For each k , we evaluate all $\binom{m}{k}$ projected designs and determine the worst projection with respect to four criteria.

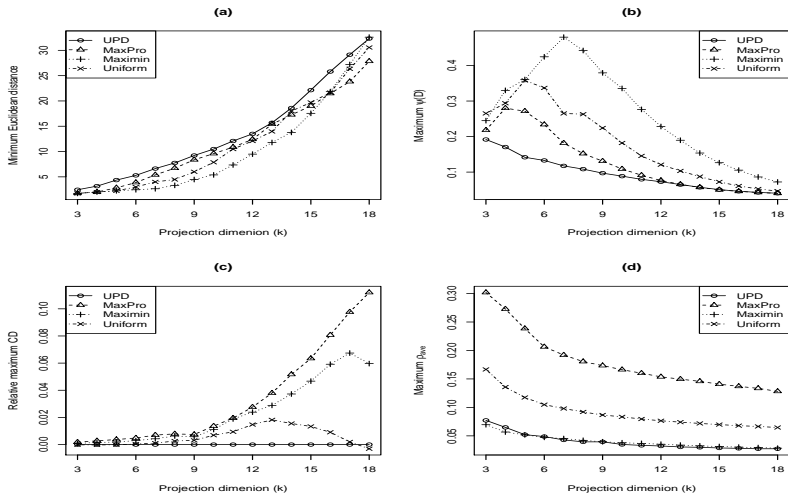


Figure 3: (a) minimum Euclidean distance (the larger the better), (b) maximum $\psi(D)$ (the smaller the better), (c) relative maximum CD (the smaller the better), and (d) maximum ρ_{ave} (the smaller the better).

Construction Methods

- Good Lattice Point (GLP) designs are LHDs and often used to construct uniform designs (Fang and Wang, 1994).
- Let $h_1 < \dots < h_p$ be p integers (from 1 to n) coprime to n

$$D = (x_{ij}) \text{ with } x_{ij} = i \times h_j \pmod{n}$$

An example $n = 7$:

$$D = \begin{pmatrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 2 & 4 & 6 & 1 & 3 & 5 \\ 3 & 6 & 2 & 5 & 1 & 4 \\ 4 & 1 & 5 & 2 & 6 & 3 \\ 5 & 3 & 1 & 6 & 4 & 2 \\ 6 & 5 & 4 & 3 & 2 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \quad \begin{array}{l} \text{with } d(D) = 12 \\ \text{(while } d_{upper} = 16) \end{array}$$

Results from Zhou and Xu (2015, Biometrika)

- An upper bound (for L_1 -distance): For any $N \times n$ LHD D ,

$$d(D) \leq d_{upper} = \lfloor (N + 1)n/3 \rfloor,$$

where $\lfloor x \rfloor$ is the integer part of x .

- Obtain the distances for four classes of GLP designs.
 - For an odd prime n , the $n \times (n - 1)$ GLP design has $d(D) = (n + 1)(n - 1)/4$.
 - the upper bound is $d_{upper} = (n + 1)(n - 1)/3$
- The $d_{eff}(D) = d(D)/d_{upper}$ for a GLP design is 75%.
- A surprising result: any **linear** level permutation of any column **does not decrease** the distance $d(D)$.

GLP + Linear Permutation

Example: $n = 7$: Total $7^6 = 117,649$ linear permutations.

- consider only 7 simple permutations: $D_i = D + i \pmod n$

$$D \quad \rightarrow \quad D_1 = D + 1 \pmod n$$

$$\begin{pmatrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 2 & 4 & 6 & 1 & 3 & 5 \\ 3 & 6 & 2 & 5 & 1 & 4 \\ 4 & 1 & 5 & 2 & 6 & 3 \\ 5 & 3 & 1 & 6 & 4 & 2 \\ 6 & 5 & 4 & 3 & 2 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \rightarrow \begin{pmatrix} 2 & 3 & 4 & 5 & 6 & 0 \\ 3 & 5 & 0 & 2 & 4 & 6 \\ 4 & 0 & 3 & 6 & 2 & 5 \\ 5 & 2 & 6 & 3 & 0 & 4 \\ 6 & 4 & 2 & 0 & 5 & 3 \\ 0 & 6 & 5 & 4 & 3 & 2 \\ 1 & 1 & 1 & 1 & 1 & 1 \end{pmatrix}$$

$$d(D) = 12$$

$$d(D_1) = 13$$

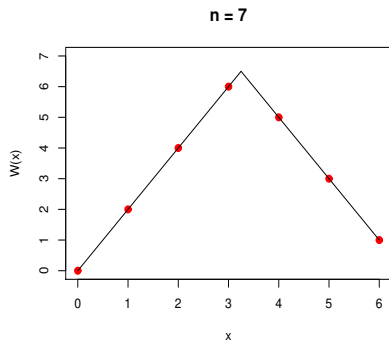
- After linear permutations, d_{eff} is about 90% for large n .

How About Nonlinear Permutations?

- Given an integer n , for $x = 0, \dots, n - 1$,

$$W(x) = \begin{cases} 2x, & \text{for } 0 \leq x < n/2; \\ 2(n - x) - 1, & \text{for } n/2 \leq x < n. \end{cases}$$

- The W is a permutation of $\{0, \dots, n - 1\}$.



The W has been useful in

1. Latin squares, Williams (1949)
2. Orthogonal designs, Bailey (1982), Edmondson (1993)
3. Orthogonal LHDs under a second-order Fourier model, Butler (2001)

GLP + Williams Transformation

Algorithm (Wang, Xiao and Xu, 2018, Annals of Statistics)

Step 1. Generate an $n \times p$ GLP design D .

Step 2. For $b = 0, \dots, n - 1$, generate $D_b = D + b \pmod{n}$.

Step 3. Let $E_b = W(D_b)$.

Step 4. Find the best E_b which maximizes $d(E_b)$.

Example: $n = 7, b = 1$

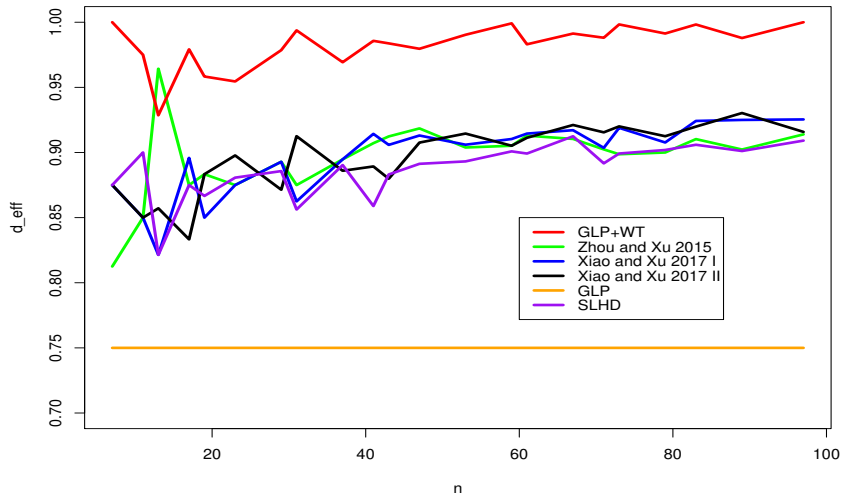
$$D \quad \rightarrow \quad D_1 = D + 1 \pmod{n} \quad \rightarrow \quad E_1 = W(D_1)$$
$$\begin{pmatrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 2 & 4 & 6 & 1 & 3 & 5 \\ 3 & 6 & 2 & 5 & 1 & 4 \\ 4 & 1 & 5 & 2 & 6 & 3 \\ 5 & 3 & 1 & 6 & 4 & 2 \\ 6 & 5 & 4 & 3 & 2 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \rightarrow \begin{pmatrix} 2 & 3 & 4 & 5 & 6 & 0 \\ 3 & 5 & 0 & 2 & 4 & 6 \\ 4 & 0 & 3 & 6 & 2 & 5 \\ 5 & 2 & 6 & 3 & 0 & 4 \\ 6 & 4 & 2 & 0 & 5 & 3 \\ 0 & 6 & 5 & 4 & 3 & 2 \\ 1 & 1 & 1 & 1 & 1 & 1 \end{pmatrix} \rightarrow \begin{pmatrix} 4 & 6 & 5 & 3 & 1 & 0 \\ 6 & 3 & 0 & 4 & 5 & 1 \\ 5 & 0 & 6 & 1 & 4 & 3 \\ 3 & 4 & 1 & 6 & 0 & 5 \\ 1 & 5 & 4 & 0 & 3 & 6 \\ 0 & 1 & 3 & 5 & 6 & 4 \\ 2 & 2 & 2 & 2 & 2 & 2 \end{pmatrix}$$

$$d(D) = 12$$

$$d(D_1) = 13$$

$$d(E_1) = 16 (= d_{upper})$$

Comparison of Various $n \times (n - 1)$ LHDs



Key Result

Let

$$b = W^{-1} \left(\frac{n-1}{2} \pm c \right)$$

where $c = \lfloor \sqrt{(n^2 - 1)/12} \rfloor$.

Theorem 3

Given a prime n and $p = n - 1$, such defined b leads the best E_b , with

$$d_{\text{eff}}(E_b) \geq 1 - 2/\sqrt{3(n^2 - 1)}.$$

As $n \rightarrow \infty$, $d_{\text{eff}}(E_b) \rightarrow 1$.

- No need for computer search: $D \rightarrow D_b \rightarrow E_b = W(D_b)$
- Guaranteed efficiency
- Larger n , better design

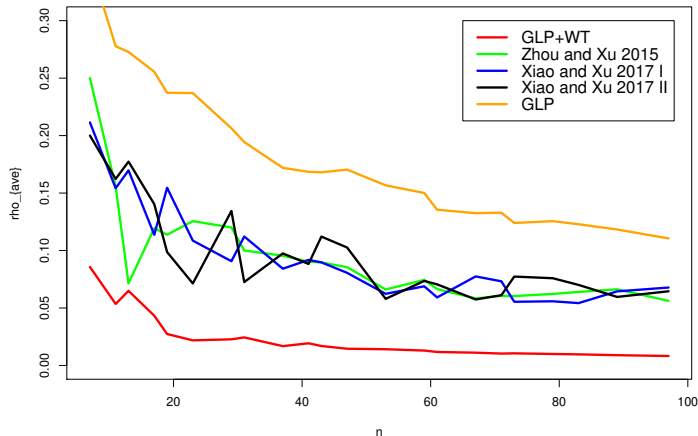
Correlations

For any $n \times p$ design $D = (x_{ij})$, define

$$\rho_{ave}(D) = \frac{\sum_{j \neq k} |\rho_{jk}|}{p(p-1)},$$

where ρ_{jk} is the correlation between columns j and k of D .

Comparison of ρ_{ave} for $n \times (n - 1)$ Designs



Method I: $\rho_{ave}(E_b) < 2/(n - 2)$

Summary

- Many available algorithms for constructing space-filling designs, but not efficient for constructing large designs
- A breakthrough — Wang, Xiao and Xu (2018) constructed maximin distance designs via good lattice points and a nonlinear transformation **without computer search**.
 - Large distance efficiencies: $d_{eff} = 1$ (or $\rightarrow 1$)
 - Low average correlation $\rho_{ave} \rightarrow 0$ as $n \rightarrow \infty$
 - Asymptotically optimal under the uniform projection criterion.
- A new class of space-filling designs — Uniform projection designs
 - suitable when only a subset of the input variables are active.
 - good space-filling not only in two dimensions, but also in all dimensions.
 - equivalent to maximin L_1 -distance criterion if L_1 -equidistant designs exist.
 - robust under other design criteria.
- There are still many open problems to be investigated.

Selected References

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