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Design and Analysis of Complex Computer Models

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Abstract

This chapter presents a review of some state-of-theart statistical techniques for analyzing real computer experiments which play a significant role in various scientific research and industrial applications. In computer experiments, emulators (i.e. surrogate models) are often used to rapidly approximate the outcomes and reduce the computational expense. Gaussian process (GP) models, also known as Kriging, are a common choice of emulators, and optimal experimental designs should be used to improve their accuracy. Specifically, space-filling designs are widely used in the literature, which proved to be efficient under GP models. In this chapter, we review different types of GP models as well as various kinds of space-filling designs. We further provide a practical tutorial on how to construct space-filling designs and fit GP emulators to analyze real computer experiments.

Keywords

Computer experiments

Gaussian process models

Space-filling designs

Latin hypercube designs

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Design and Analysis of Complex Computer Models



Jeevan Jankar, Hongzhi Wang, Lauren Rose Wilkes, Qian Xiao, and Abhyuday Mandal

Abstract This chapter presents a review of some state-of-the-art statistical tech-1 niques for analyzing real computer experiments which play a significant role in 2 various scientific research and industrial applications. In computer experiments, 3 emulators (i.e. surrogate models) are often used to rapidly approximate the out-Δ comes and reduce the computational expense. Gaussian process (GP) models, also 5 known as Kriging, are a common choice of emulators, and optimal experimental 6 designs should be used to improve their accuracy. Specifically, space-filling designs 7 are widely used in the literature, which proved to be efficient under GP models. 8 In this chapter, we review different types of GP models as well as various kinds 9 of space-filling designs. We further provide a practical tutorial on how to construct 10

space-filling designs and fit GP emulators to analyze real computer experiments.

¹² **Keywords** Computer experiments · Gaussian process models · Space-filling

designs · Latin hypercube designs

14 **1 Introduction**

A computer experiment is a system of complex computer codes simulating a phys-15 ical process. They are implemented like a function, taking inputs to produce the 16 outputs. This automation can reduce the cost, time, and/or management compared 17 to a traditional lab experiment (see, for example, [20]). Computer experiments are 18 often deterministic (specified inputs will always produce the same output), making 19 the results more stable and less prone to random errors compared to traditional lab 20 experiments. Researchers can manipulate the code to systematically adjust a wide 21 range of inputs and generate outputs based on what they are trying to study. They 22 are instrumental in cases where a physical experiment would be impossible, such 23 as modeling black holes [29]. Due to these characteristics, computer experiments 24

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²⁵ become very popular in various scientific research and industrial applications (see,
²⁶ for more examples, [12, 20]). For example, [8] created a 3D mixed finite element
²⁷ model to study flexoelectric material. The Flexoelectric Effect is where strain gra²⁸ dients polarize electric fields. This process is complicated to study, especially in a
²⁹ practical context, so the finite element method is a numerical approach, i.e. computer
³⁰ experiment, used to study this effect. Mixed finite elements simplify this task further
³¹ using an alternative way of handling higher order derivatives.

Computer experiments are often computationally intensive, though computing 32 power has increased in recent years. To rapidly generate many outcomes and reduce 33 the computational expenses, emulators (i.e. surrogate models) are needed which are 34 often fitted with only a few data points. Emulators should also allow uncertainty 35 quantification to measure how accurate the model is for predictions. If a good emu-36 lator is selected, it may be more useful than the underlying physical process as it 37 eliminates noise. The Gaussian Process (GP) model is a widely used emulator [20, 38 43]. The GP assumes all observations following a multivariate normal distribution, 39 which is characterized by a mean vector μ and a covariance matrix Σ . The GP model 40 would interpolate the observations, which is desirable for computer experiments hav-41 ing deterministic outputs. It also allows for accurate uncertainty quantification. By 42 specifying different types of covariance functions, researchers may further add prior 43 knowledge about the shape of the response surface. 11

The GP model has been applied to many computer experiments in Chemistry, 45 Computational Biology, Robotics and others [30]. As an illustration, it has accu-46 rately simulated the collision dynamics of complex molecules [6], the spread of 47 COVID-19 [52], flagging suspicious Internet claims [63] and autonomous learning 48 in robots [7]. Data scientists at Microsoft introduced a framework that enables the <u>1</u>0 application of GP models to data sets containing millions of data points [23]. As 50 pictured in Fig. 1, a Bayesian framework is used for human body pose tracking [10]. 51 Instead, a GP experiment can be used to take in a description of a human silhou-52 ette as inputs and outputs to identify human pose [68]. One useful application of 53 GP in Astronomy is modeling the collision of two black holes. Researchers cannot 54 create black holes to observe and experiment with, so computer experiments offer 55 a veritable way to simulate the outcome of black hole collisions. Figure 2 illustrates 56



Fig. 1 An example of Bayesian framework for human pose tracking *Source* https://www.ncbi. nlm.nih.gov/pmc/articles/PMC3292173/ [68]

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Fig. 2 Computer simulation of two black holes colliding *Source* https://www.black-holes.org/code/SpEC.html

- 57 that computer models and GP emulators are created based on the known properties
- of black holes and the surrounding system of space and are compared to naturally
- ⁵⁹ observed black hole movement in order to test how accurate they are [58]. Another
- ⁶⁰ interesting application of GP is on car crash simulation to study the damage on the
- 61 car. Here, models are validated by comparing simulation results with an actually
- ⁶² controlled crash. Figure 3 depicts some results from a finite element method.



Fig. 3 An example of Gaussian Process experiment in car crash simulation *Source* https://www. csm.ornl.gov/SC98/car.html

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The remainder of this chapter is organized as follows. In Sect. 2, we systematically review the GP models. Specifically, we discuss the ordinary and universal GP in Sect. 2.1, their model estimations and uncertainty quantification in Sect. 2.2 and methods for including qualitative inputs in Sect. 2.3. In Sect. 3, we review popular experimental designs used in computer experiments, and we conclude this chapter in Sect. 4.

69 2 The Gaussian Process Model

⁷⁰ In this section, we aim to understand GP as a flexible nonparametric regression for ⁷¹ surrogate modeling in computer experiments. GP is widely used in many statistical ⁷² and probabilistic modeling enterprises. GP is a very generic term, and all it means ⁷³ is that any finite collection of realizations is modeled as having a multivariate nor-⁷⁴ mal (MVN) distribution. That means, a finite collection of *n* observations can be ⁷⁵ completely characterized by their mean vector μ and covariance matrix Σ . ⁷⁶ Let $y(\mathbf{x_i})$ be the output which is assumed to be a deterministic real-valued function

of the *d*-dimensional variable $\mathbf{x_i} = (x_{i1}, \dots, x_{id})^T \in D \subset \mathbb{R}^d$, for $i = 1, 2, \dots, n$. Let $(Y_x)_{x \in D}$ be a square-integrable random field and y be a realization of $(Y_x)_{x \in D}$. Let $\mathbf{X} = \{\mathbf{x_1}, \dots, \mathbf{x_n}\}$ be the points where their responses have been observed, which

is denoted by $\mathbf{y} = (y(\mathbf{x_1}), \dots, y(\mathbf{x_n}))^T$. The aim of GP is to optimally predict Y_x by a linear combination of the observations \mathbf{y} , for any $\mathbf{x} \in D$.

82 2.1 Model Formulation

⁸³ Ordinary GP, also known as ordinary Kriging, has the form

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$$y(\mathbf{x}_{\mathbf{i}}) = \mu + Z(\mathbf{x}_{\mathbf{i}}),\tag{1}$$

where μ is the mean vector and $Z(\mathbf{x}_i)$ is a GP such that $Z(\mathbf{x}_i) \sim GP(0, \sigma^2 \Sigma)$. 85 In the above model, $Z(\mathbf{x}_i)$ is GP with zero mean, and the covariance function 86 $\phi(\cdot) = \sigma^2 \Sigma(\cdot|\theta)$, where $\theta = (\theta_1, \dots, \theta_d)^T$ is the vector of unknown correlation 87 parameters with all $\theta_k > 0$ (k = 1, ..., d) and Σ is a stationary correlation func-88 tion that determines the correlation between inputs with parameters `. The mean of 89 the GP controls the trend, whereas the correlation function controls the smoothness 90 of its sample paths. Power-exponential, Gaussian and Matérn correlation functions 91 are the most widely used ones in the literature. 92

In the power-exponential correlation structure, the (i, j)th element in the correlation matrix is defined as follows: Design and Analysis of Complex Computer Models

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Author Proof

$$\Sigma\left(\mathbf{x}_{i}, \mathbf{x}_{j} \mid \boldsymbol{\theta}\right) = \prod_{k=1}^{d} \exp\left\{-\theta_{k} \left|x_{ik} - x_{jk}\right|^{p_{k}}\right\} \quad \text{for all } i, j,$$
(2)

with two inputs $\mathbf{x}_i = (x_{i1}, \ldots, x_{id})^T$ and $\mathbf{x}_j = (x_{j1}, \ldots, x_{jd})^T$ and smoothness parameters p_1, \ldots, p_d which lie between 0 and 2, with 0 giving the most rough results and 2 giving the most smooth. If we take $p_k = 2$ for all $k = 1, \ldots, d$, then it results in the popular Gaussian correlation function:

$$\Sigma\left(\mathbf{x}_{i}, \mathbf{x}_{j} \mid \boldsymbol{\theta}\right) = \prod_{k=1}^{d} \exp\left\{-\theta_{k} \left|x_{ik} - x_{jk}\right|^{2}\right\} \quad \text{for all } i, j.$$
(3)

¹⁰¹ The correlation functions of Matérn family is given by

$$\Sigma(\mathbf{h} \mid \boldsymbol{\theta}) = \prod_{k=1}^{d} \frac{1}{\Gamma(v)2^{v-1}} \left(\frac{2\sqrt{v} \mid h_k \mid}{\theta_k}\right)^v K_v \left(\frac{2\sqrt{v} \mid h_k \mid}{\theta_k}\right), \tag{4}$$

where v > 0 is a smoothness parameter, $\Gamma(\cdot)$ is the Gamma function and $K_v(\cdot)$ is the modified Bessel function of order v. Two commonly used orders are v = 3/2 and v = 5/2.

Different correlation functions mentioned above impose different characteristics for function draws, allowing for different properties when modeling computer models. For example, when using the power-exponential function, all sample paths are infinitely differentiable when $p_k = 2$. For the Matérn correlation function, when we have d = 1, all sample paths are $\lceil v \rceil - 1$ differentiable. Hence, v is viewed as a smoothness parameter.

In the literature, two important assumptions are often imposed on the ordinary GP 112 model to effectively analyze computer experiment. One assumption is that the GP 113 is separable [9], which means finite-dimensional distributions can determine sample 114 path properties of function draws which are usually infinite-dimensional. The sec-115 ond important assumption is that the model is stationary. Consider $\{x_1, \ldots, x_n\} \in D$ 116 and any $h \in \Re^d$, then a GP model is said to be stationary if the random vectors 117 $(Y(\mathbf{x_1}), \ldots, Y(\mathbf{x_n}))$ and $(Y(\mathbf{x_1} + \mathbf{h}), \ldots, Y(\mathbf{x_n} + \mathbf{h}))$ follow the same distribution. 118 This means that both these random vectors should have the same mean and covari-119 ance. 120

The second assumption is restrictive, and we may need more flexibility while modeling computer experiments. One popular approach is to extend the above ordinary GP model to incorporate a global trend function for the mean part. This extended model is known as *Universal Kriging* which has the form:

$$\mathbf{y}(\mathbf{x}) = \mu(\mathbf{x}) + Z(\mathbf{x}),\tag{5}$$

with $\mu(\mathbf{x}) = \mathbf{f}(\mathbf{x})^T \boldsymbol{\beta} = \sum_{s=1}^m \beta_s f_s(\mathbf{x})$, where **f** is a *m*-dimensional known function and $\boldsymbol{\beta} = (\beta_1, \dots, \beta_m)^T$ is a vector of unknown parameters. The idea is to rely on **Author Proof**

functions in $\mathbf{f}(\mathbf{x})$ to de-trend the process and then model any residual variation as zero mean stationary GP. Taking constant mean $\mathbf{f}(\mathbf{x}) \equiv 1$ results in the ordinary GP model discussed above. The stationary correlation functions discussed above in Eqs. (2) and (4) can also be applied here, that is,

$$Cov\left(Z(\mathbf{x}+\mathbf{h}), Z(\mathbf{x})\right) = \sigma^2 \Sigma(\mathbf{h}),$$

where correlation function $\Sigma(\mathbf{h})$ is a positive semidefinite function with $\Sigma(\mathbf{0}) = 1$ and $\Sigma(\mathbf{h}) = \Sigma(-\mathbf{h})$.

135 2.2 Estimation and Uncertainty Quantification

In this section, we present equations used for predicting and quantifying uncertainty on $y(\mathbf{x})$ given observed responses $\mathbf{y} = (y(\mathbf{x_1}), \dots, y(\mathbf{x_n}))^T$. The question we are trying to answer is: given examples of function in pairs $(\mathbf{x_1}, y(\mathbf{x_1})), \dots, (\mathbf{x_n}, y(\mathbf{x_n}))$, what random function realizations could explain or could have generated those observed values? In other words, we want to calculate the conditional distribution $(Y(\mathbf{x_1}), \dots, Y(\mathbf{x_n})) |\{(\mathbf{x_1}, y(\mathbf{x_1})), \dots, (\mathbf{x_n}, y(\mathbf{x_n}))\}.$

Before we calculate the *predictive distribution*, we need to address the key question of how the parameters β , σ^2 and θ are estimated from data $(\mathbf{x}_i, y(\mathbf{x}_i))_{i=1}^n$. The most popular approach for parameter estimation is *maximum likelihood estimation*, and the log-likelihood function under the above assumed GP model can be written as

¹⁴⁶
$$l\left(\boldsymbol{\beta},\sigma^{2},\boldsymbol{\theta}\right) = -\frac{1}{2}\left[n\log\sigma^{2} + \log\det\Sigma_{\boldsymbol{\theta}} + \frac{1}{\sigma^{2}}\left(\mathbf{y} - \mathbf{F}\boldsymbol{\beta}\right)\Sigma_{\boldsymbol{\theta}}^{-1}\left(\mathbf{y} - \mathbf{F}\boldsymbol{\beta}\right)\right],$$
 (6)

where det Σ_{θ} is the determinant of the matrix $\Sigma_{\theta} = \left[\Sigma(\mathbf{x}_{i}, \mathbf{x}_{j})\right]_{i=1}^{n} \sum_{j=1}^{n} \text{ and } \mathbf{F} = [f_{s}(\mathbf{x}_{i})]_{i=1}^{n} \sum_{s=1}^{m}$. Hence, the MLEs for $(\boldsymbol{\beta}, \sigma^{2}, \boldsymbol{\theta})$ are the parameter estimates that maximize the above log-likelihood function. ML estimates of $(\boldsymbol{\beta}, \sigma^{2})$ for fixed value of $\boldsymbol{\theta}$ can be easily obtained as follows:

$$\hat{\boldsymbol{\beta}}_{\boldsymbol{\theta}} = \left(\mathbf{F}^T \boldsymbol{\Sigma}_{\boldsymbol{\theta}}^{-1} \mathbf{F}\right)^{-1} \mathbf{F}^T \boldsymbol{\Sigma}_{\boldsymbol{\theta}}^{-1} \mathbf{y}$$
(7)

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$$\hat{\sigma}_{\theta}^{2} = \frac{1}{n} \left(\mathbf{y} - \mathbf{F} \hat{\boldsymbol{\beta}}_{\theta} \right)^{T} \boldsymbol{\Sigma}_{\theta}^{-1} \left(\mathbf{y} - \mathbf{F} \hat{\boldsymbol{\beta}}_{\theta} \right).$$
(8)

Substituting these ML estimates back into Eq. (6), we get the profile likelihood function as follows:

 $l\left(\hat{\boldsymbol{\beta}}, \hat{\sigma}^2, \boldsymbol{\theta}\right) = -\frac{1}{2} \left[n \log \hat{\sigma}^2 + \log \det \Sigma_{\boldsymbol{\theta}} + n \right], \tag{9}$

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where the MLE of θ is one that maximizes the above function in Eq. (9). This optimization problem does not enjoy a closed-form solution, and numerical methods, e.g. quasi-Newton algorithms [40] are used for solving the problem.

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Once we have estimates of parameters, we can calculate the conditional distribution as mentioned above. Let $(\hat{\beta}, \hat{\sigma}^2, \hat{\theta})$ denote the ML estimates of unknown parameters for the given GP model. Then for a new input $\mathbf{x}^* \in \mathbb{R}^d$, the mean and variance of random variable $Y(\mathbf{x}^*|\mathbf{y})$ are as follows:

$$\hat{y}\left(\mathbf{x}^{*}\right) = \mathbb{E}\left[Y\left(\mathbf{x}^{*}\right) \mid \mathbf{y}\right] = \mathbf{f}^{T}\left(\mathbf{x}^{*}\right)\hat{\boldsymbol{\beta}} + \mathbf{r}_{\hat{\theta}}^{T}\left(\mathbf{x}^{*}\right)\boldsymbol{\Sigma}_{\hat{\theta}}^{-1}\left(\mathbf{y} - \mathbf{F}\hat{\boldsymbol{\beta}}\right),$$
(10)

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$$s\left(\mathbf{x}^{*}\right)^{2} = \operatorname{Var}\left[Y\left(\mathbf{x}^{*}\right) \mid \mathbf{y}\right] = \hat{\sigma}^{2}\left(1 - \mathbf{r}_{\hat{\theta}}^{T}\left(\mathbf{x}^{*}\right)\Sigma_{\hat{\theta}}^{-1}\mathbf{r}_{\hat{\theta}}\left(\mathbf{x}^{*}\right)\right), \quad (11)$$

where the covariance vector $\mathbf{r}_{\hat{\theta}}(\mathbf{x}^*) = \left[\Sigma_{\hat{\theta}}(\mathbf{x}^*, \mathbf{x}_1), \Sigma_{\hat{\theta}}(\mathbf{x}^*, \mathbf{x}_2), \dots, \Sigma_{\hat{\theta}}(\mathbf{x}^*, \mathbf{x}_n) \right]^{\mathrm{T}}$. 167 When some observed data points are very close to each other, the covariance 168 matrix $\Sigma_{\hat{\theta}}$ may become nearly singular, making it difficult to obtain a stable inverse 169 matrix Σ_{a}^{-1} . This is a common issue for GP models, when the run and/or factor sizes 170 are large. One way to deal with this problem is to add a positive scalar λ , called the 171 *nugget* parameter, to the diagonal elements in $\Sigma_{\hat{\theta}}$, i.e. replacing Σ_{θ} with $\Sigma_{\theta} + \lambda \mathbf{I}$, 172 where I is an identity matrix. Adding λ is analogous to adding the ridge parameter 173 in ridge regression, which helps in moving the smallest eigenvalue of Σ_{θ} away from 174 zero, thus stabilizing the calculation of its inverse. 175

For large data sizes, the estimation of GP models can be very time-consuming, 176 mainly due to the matrix inverse calculation of order $O(n^3)$. To deal with this problem, 177 [21] proposed a localize GP (LaGP) approach. Based on a local subset of the data, they 178 provide a family of local sequential design schemes that defines the support points 179 of a GP predictor. The idea is to make sure that for a given choice of covariance 180 structure, the data points far from the target location \mathbf{x}^* will have little effect on 181 the prediction. Hence, it is not unwise to calculate the inverse of the full covariance 182 matrix, as the elements corresponding to "far away" points will contribute very little 183 to predicting $y(\mathbf{x}^*)$. Interested readers may refer to [21] for further details. 184

The notion of calibration and sensitivity analysis is important in the context of 185 physical and computer experiments. In practice, we only observe response y_{Field} 186 instead of observing real physical response y_{Real} . And, we use the above computer 187 models to approximate y_{Real} as y_{Model} . Now, as we saw in the earlier sections apart 188 from input variables, computer models also use some more parameters known as 189 calibration parameters to fine-tune the model. Covariance parameters θ are one such 190 example of calibration parameters. A Bayesian framework was proposed by [28] to 191 address this as follows: 192

$$y_{Real}(\mathbf{x}) = y_{Model}(\mathbf{x}, \boldsymbol{\theta}) + \mathbf{b}(\mathbf{x})$$

$$y_{Field}(\mathbf{x}) = y_{Model}(\mathbf{x}, \boldsymbol{\theta}) + \mathbf{b}(\mathbf{x}) + \epsilon,$$

where $b(\mathbf{x})$ is a bias and ϵ is the normal error. Reference [28] used Bayesian methods to estimate the bias correction function and unknown calibration parameter θ under a GP prior. Iterative history matching algorithm as one proposed by [53] for calibrating a galaxy formation model called GALFORM is an alternative to this Bayesian approach. Recently, [1] used this algorithm for calibrating hydrological time-series models.

201 2.3 GP with Qualitative Inputs

The above-mentioned GP model is valid only with quantitative inputs, but there are 202 many situations in real life where inputs can be both quantitative and qualitative. 203 One straightforward way to adapt GP models with qualitative inputs is to construct 204 separate GP models for each level combination of the qualitative factors. Yet, when 205 there are many high-level qualitative factors, such an approach would require many 206 observations to fit a large number of GP models. In the current literature, many 207 integrated GP models for both quantitative and qualitative factors are proposed [22, 208 41, 50, 65, 66]. 209

Reference [60] proposed a new method called EzGP to deal with such problems. Let the *k*th input of the computer emulator be $\mathbf{w}_k = (\mathbf{x}_k^T, \mathbf{z}_k^T)^T$, where $\mathbf{x}_k = (x_{k1}, \ldots, x_{kp})^T$ is the continuous part of input as mentioned in the previous sections and $\mathbf{z}_k = (z_{k1}, \ldots, z_{kq})^T \in \mathbb{N}^q$ is the qualitative part of the input, where $k = 1, \ldots, n$. The EzGP method is inspired by the idea of Analysis of Variance (ANOVA) where quantitative and qualitative inputs are jointly modeled as follows:

$$y(\mathbf{w}) = \mu + Z_{\mathbf{z}}(\mathbf{x}), \tag{12}$$

which suggests that for any given level combination of qualitative factors, $y(\mathbf{w})$ is a GP. Specifically, they considered the following additive model structure:

$$Z_{\mathbf{z}}(\mathbf{x}) = Z_0(\mathbf{x}) + Z_{z^{(1)}}(\mathbf{x}) + \dots + Z_{z^{(q)}}(\mathbf{x}),$$
(13)

where Z_0 and $Z_{\tau^{(h)}}$ for $h = 1, \dots, q$ are independent GPs with mean zero and some 220 covariance functions. Here, Z_0 plays the role of base GP which takes only quantitative 221 inputs reflecting the intrinsic relation between y and x, and other GPs $Z_{z^{(h)}}$ are the 222 adjustments made to the base GP to reflect the impact of each qualitative factor $z^{(h)}$ 223 for $h = 1, \ldots, q$. The EzGP method can easily deal with heterogeneity in computer 224 models with multiple qualitative factors. Two variants in EzGP are proposed to fit data 225 with high dimensionality or large run sizes, which can achieve high computational 226 efficiency. 227

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3 Designs for Computer Experiments

²²⁹ Computer codes generate outputs in a deterministic manner in computer experiments, ²³⁰ meaning the same input returns the same output (no random errors). Latin hypercube ²³¹ designs (LHDs, [38]) are the most popular experimental designs in computer exper-²³² iments. An *n* runs and *k* factors, LHD is an $n \times k$ matrix with each column being a ²³³ random permutation of numbers 1, ..., *n*. LHDs do not have replicates in each one-²³⁴ dimensional projection. There are various types of optimal LHDs for practical needs, ²³⁵ including space-filling LHDs, maximum projection LHDs and orthogonal LHDs.

When we have little or no information about the response surface, it is desirable 236 to have design points as scattered out as possible in the design space for better explo-237 ration. Despite LHDs having a uniform one-dimensional projection property, random 238 LHDs may have poor space-filling properties over the entire design space. Figure 4 239 is an illustrative example with two LHD designs. The LHD in the left panel is con-240 centrated almost entirely on the diagonal, which clearly does not explore the input 241 space sufficiently. The design points in the right panel are scattered out over the entire 242 design space, so this design may provide more reliable information. The maximin 243 distance criterion [25] is a widely used metric for measuring the space-filling prop-244 erty of LHDs. It aims to maximize the minimum distances between design points. 245 Let **X** denote an LHD matrix, where the L_a -distance between two runs x_i and x_j of **X** 246 is given by $d_q(x_i, x_j) = \left\{ \sum_{k=1}^m |x_{ik} - x_{jk}|^q \right\}^{1/q}$, where q is an integer. Two popular 247 choices are q = 1 (i.e. the Manhattan distance) and q = 2 (i.e. the Euclidean dis-248 tance). The maximin L_a -distance design has the maximized minimum L_a -distance, 249 i.e. max min $d_a(x_i, x_j)$, where $1 \le i < j \le n$. Reference [24, 39] further proposed 250 a scalar value to evaluate the maximin distance criterion: 251



Fig. 4 Latin hypercube designs for size n = 5 and k = 2

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 $\phi_p = \left\{ \sum_{i=1}^{n-1} \sum_{j=i+1}^n d_q(x_i, x_j)^{-p} \right\}^{1/p},\tag{14}$

where *p* is a tuning parameter. As $p \to \infty$, the ϕ_p criterion in Eq. (14) is asymptotically equivalent to the Maximin distance criterion, and p = 15 is usually sufficient in practice. The LHDs that minimize the ϕ_p criterion are called the maximin distance LHDs.

In the literature, both algebraic constructions (add some references) and search algorithms (add some reference) are proposed to construct maximin distance LHDs. 258 Algebraic constructions usually require very little computational cost to generate 259 optimal LHDs, which are very attractive for large design sizes. Yet, they are only 260 available for certain design sizes. Search algorithms can generate optimal designs of 261 flexible sizes, but they often require more computation resources to identify optimal 262 LHDs. As there are $(n!)^{k-1}$ possible LHDs with *n* runs and *k* factors, search algo-263 rithms could become very costly when n and k are large. Here, we will briefly survey 264 some popular construction methods; see [55] for a survey. 265

Specifically, [56] proposed to generate maximin distance LHDs via good lat-266 tice point (GLP) sets [67] and Williams transformation [59]. They proved that the 267 resulting designs of sizes $n \times (n - 1)$ (with *n* being any odd prime) and $n \times n$ (with 268 2n + 1 or n + 1 being odd prime) are optimal under the maximin L_1 -distance crite-260 rion. The construction method starts by generating a GLP design, and then use the 270 Williams transformation [59] to improve a linear permuted GLP design. Reference 271 [51] proposed to construct orthogonal array-based LHDs (OALHDs) from existing 272 orthogonal arrays (OAs). The key idea of this construction is to deterministically 273 replace OA entries with a random permutation of LHD elements. OALHDs inherit 274 the properties of OAs and tend to have better space-filling properties compared to 275 random LHDs. Note that the design sizes of OALHDs rely on the existence of cor-276 responding OAs. 277

Search algorithms should be used to generate optimal LHDs when no construction 278 methods are available. Reference [39] proposed a simulated annealing (SA) algo-279 rithm, which randomly exchanges elements to seek improvements over iterations to 280 identify global best LHDs. Following the work of [31, 39, 51] proposed to construct AQ2281 orthogonal array-based LHDs (OALHDs) using the SA algorithm. They proposed to 282 exchange elements that share the same original OA entry randomly. Reference [27] 283 proposed a multi-objective criterion and developed a modified SA algorithm to gener-284 ate optimal LHDs having good space-filling properties as well as orthogonality. This 285 algorithm can lead to many good designs, but it is often computationally heavy, since 286 it calculates all average pairwise correlations and row-wise distances at each itera-287 tion. Besides these SA-based algorithms, [32] proposed to use a genetic algorithm 288 (GA) for searching optimal designs, which focuses on global best by exchanging 289 random columns between global best and other candidate solutions. In addition, [3] 290 proposed a version of the particle swarm optimization (PSO) algorithm by gradually 291 reducing the Hamming distances between each particle and its personal best (or the 292

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²⁹³ global best). Generally speaking, the PSO is recommended for small design sizes ²⁹⁴ $(n \le 7)$ and the GA has better performance for moderate and large design sizes.

²⁹⁵ Uniform designs (UDs) [11, 13] are another popular type of space-filling designs. ²⁹⁶ There are various measurements of uniformity proposed in the literature, such as ²⁹⁷ the star L_2 -discrepancy [57], modified L_2 -discrepancy [14] and the centered L_2 -²⁹⁸ discrepancy [15]. The search algorithms mentioned above can be used for identifying ²⁹⁹ UDs.

Maximin distance LHDs have space-filling properties in the full-dimensional space, but their two to k - 1-dimensional projections may not be space-filling. Reference [26] proposed the maximum projection LHDs (MaxPro LHDs) which enhance the space-filling properties in all possible dimensional projections. Analogous to [26, 39] defined the maximum projection criterion as

$$\min_{\mathbf{X}} \psi(\mathbf{X}) = \left\{ \frac{1}{\binom{n}{2}} \sum_{i=1}^{n-1} \sum_{j=i+1}^{n} \frac{1}{\prod_{l=1}^{k} (x_{il} - x_{jl})^2} \right\}^{1/k}.$$
 (15)

³⁰⁶ LHDs that minimize the ψ values are called MaxPro LHDs. Reference [26] pro-³⁰⁷ posed an SA-based search algorithm to identify MaxPro LHDs.

Orthogonal LHDs (OLHDs) are another type of optimal LHDs which aim to minimize the correlations between factors [16, 45, 48]. Two correlation-based criteria are often used to measure designs' orthogonality: the average absolute correlation criterion and the maximum absolute correlation criterion [16], which are defined as

ave
$$(|q|) = \frac{2\sum_{i=1}^{k-1}\sum_{j=i+1}^{k} |q_{ij}|}{k(k-1)}$$
 and $\max|q| = \max_{i,j} |q_{ij}|,$ (16)

where q_{ij} is the correlation between the *i*th and *j*th columns in the design matrix. Orthogonal designs may not exist for all sizes. In practice, designs with small ave(|q|) or max|q| are preferred.

In the literature, construction methods of OLHDs are widely explored. Specifi-316 cally, [62] proposed a method to construct OLHDs with run sizes $n = 2^m + 1$ and 317 factor sizes k = 2m - 2, where m is any integer no less than 2. Reference [5] extended 318 the work of [62] to accommodate more factors. Reference [45] developed a method 319 based on factorial designs with group rotations for $n = 2^{2^m}$ and $k = 2^m t$, where 320 m is any positive integer and t is the number of rotation groups. Reference [47] 321 improved their earlier work [46] to construct OLHDs with even more flexible run 322 sizes: $n = r2^{c+1}$ or $n = r2^{c+1} + 1$ and $k = 2^c$, where c and r are any two positive 323 integers. Reference [61] proposed to use generalized orthogonal designs to construct 324 OLHDs and nearly orthogonal LHDs (NOLHDs) with $n = 2^{r+1}$ or $n = 2^{r+1} + 1$ 325 and $k = 2^r$, where r is any positive integer. Reference [17] proposed to take advan-326 tage of OAs and their full fold-overs for OLHDs with n = 2ak runs and k factors, 327 where k is the size of orthogonal matrix and a is any positive integer. Reference [2] 328 implemented the Williams transformation [59] to construct OLHDs with odd prime 329 run-size *n* and factor-size $k \le n - 1$. Reference [33] proposed to couple OLHDs or

AQ3

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NOLHDs with OAs to accommodate large numbers of factors with fewer runs: n^2 runs and 2fp factors, where *n* and *p* are design sizes of the OLHDs or NOLHDs and 2f is the number of columns in the coupled OA.

333 4 Discussion

There are many instances in nature where it is either expensive or impossible to 334 conduct a physical experiment. For example, it is prohibitively difficult to conduct a 335 study for investigating the devastation caused by a nuclear explosion. Instances like 336 the formation of a galaxy or the formation of binary black holes cannot be studied 337 through physical experiments. Computer experiments can simulate such phenom-338 ena with reasonable accuracy. Although such computer simulators are a lot more 339 desirable than real experiments, they are still computationally expensive. To deal 340 with this problem, scientists use surrogates (emulators) to facilitate the analysis and 341 optimization of complex systems. GPs are widely used as surrogates (or emulators). 342 Space-filling designs, such as LHDs, are often used to reap the benefits of utilizing 343 such surrogates effectively. 344

Several efficient packages in R are available for fitting the GP model and identifying LHDs. Interested readers can explore different packages for fitting GP: Local
Approximate Gaussian Process Regression (laGP) by [19], DiceKriging (Kriging
Methods for Computer Experiments) by [42] and GP-fit (Gaussian Processes Modeling) by [36]. For obtaining LHDs with flexible run sizes, packages like Latin
Hypercube Designs (LHD) by [54] and Maximin-Distance (Sliced) Latin Hypercube Designs (SLHD) by [44] can be used.

Even though the computing power has increased dramatically over the last few 352 years, handling big data remains a challenging problem. There is an increasing body 353 of literature for computer experiments with large numbers of data points, but the 354 existing literature on large numbers of input variables is still meager. For details, 355 please refer to the review article by [35]. The problem of data reduction is an active 356 area of research among statisticians and computer scientists, and much progress 357 needs to be done in this area. Recent work on this includes techniques like kernel 358 handling [4] and support points [37]. 359

Different Bayesian approaches for analyzing computer experiments have been 360 discussed in the literature, particularly in the context of uncertainty quantification, 361 but most of them are difficult to implement and time-consuming [18, 28]. To solve 362 this problem, we need more advanced techniques. Another topic of active research 363 is to incorporate qualitative input variables. Many practical applications have both 364 quantitative and qualitative inputs, e.g. the data center computer experiment [41] 365 and the study of high-performance computing systems [64]. However, traditional 366 GP modeling is designated for only quantitative inputs, since its covariance function 367 of responses is constructed under the continuous input space with proper distance 368 metrics. More effective techniques and algorithms need to be developed that can 369 accommodate qualitative inputs and one such recent work is [60]. 370

Finally, there is vast existing literature on continuous response, but there are many instances where the response is binary or non-continuous. For example, binary black hole formation [34] or computer experiments with binary time series have non-Gaussian observations [49]. For handling high-dimensional input parameter space, input variables with non-continuous characteristics and non-Gaussian observations, new techniques and algorithms need to be developed.

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