Traditional or Adaptive Experimental Design? A Comparison of Statistical Design of Experiments and Bayesian Optimization for a Chemical Synthesis Problem

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Introduction





- Although designs are static, several stages are reccomended for DOE
- Classic designs tipically assumes the system can be described by a first and second order polynomial model



1. Introduction

- Recently, Bayesian Optimization (BO) has attracted attention as an alternative approach for experimental design
- BO is a sequential Bayesian experimental design framework with two main components:
 - Bayesian surrogate model that reflects **prior beliefs** and provides uncertainty estimates given data
 - Acquisition function balances **exploration** with **exploitation** to search for the optimum



- Despite recent popularity, the principles of BO are not new
 - Thompson sampling was suggested in 1930's as an heuristic to solve the exploration-exploitation dillema
 - BO was formalized in the 1960's and had a recent resurgence in the context of hyperparameter tuning

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1. Introduction

• We aim to compare sequential DOE with BO for a complex real-world chemical synthesis problem







Design focus: reducing **optimum uncertainty** Dow

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Case study

2. Case study

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- Openly available chemical reaction dataset used as case study [1]
- Problem has 3 categorical factors, 2 continuous factors and a single response (reaction yield)

12 ligands

2 continuous variables: Temperature and 4 bases 4 solvents concentration PPhMe₂ PPht-Bu₂ PCy_a GorlosPhos 120 0.057 0.153 KOAc CsOAc **BuOAc** BuCN P(t-Bu)(Ph) DMAc **KOPiv** CsOPiv Me₂N NMe p-Xylene 105 0.100 T (°C) C (M) t-BuPh-CPhos **BrettPhos** CaMe-PPh XPhos PPh₂Me PPh. P(Fur)₃ JackiePhos

- Experimental data from full factorial design (1728 samples) used as proxy for reaction system
- Objective: find the combinations of the factors that maximize reaction yield

[1] Shields et. al, 2021, Bayesian reaction optimization as a tool for chemical synthesis, Nature

2. Case study



- One-hot-encoding (OHE) is the standard approach to model categorical factors
- In chemical synthesis, we can also use chemical descriptors as covariates for each individual categorical factor
- 291 covariates are obtained from Density Functional Theory (DFT) simulations
 - Due to high dimensionality, descriptors cannot be used within standard DOE
 - BO is used with both types of encodings (OHE and DFT)

One-hot-encoding (OHE)

Base	Level 1	Level 2	Level 3
1	1	0	0
2	0	1	0
3	0	0	1

Chemical descriptors

Base	Descriptor 1	Descriptor 2	Descriptor 3	Descriptor 4
1	10	30	100	30
2	7	2	30	2
3	2	0,1	11	0,1

Categorical levels are mapped to 291 molecular features



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D-optimal screening design (main effects only)

Fit ordinary least squares model and select important factors

I-optimal design for full quadratic model

Estimate full quadratic model using LASSO (due to high number of parameters)

Find optimum factors that maximize yield

- There are 192 total possible combinations for categorical factors
- D-optimal design for main effects requires 48 experiments
- Factors are selected according to p-values and magnitude of regression coeficients
- I-optimal design is used to obtain minimum prediction variance over design space
- JMP-PRO 18 used for design and modeling



• Gaussian processes (GP) allow to obtain a **posterior distribution over functions**

 $Posterior = \frac{Likelihood \ x \ Prior}{Marginal \ Likelihood}$

- GP mean and kernel encode the prior belief about the smoothness and overall shape of the function
- GPs provide uncertainty estimates and can flexibly model both midly and highly non-linear functions



• The kernel lengthscale controls the complexity of the response surface given the distance between points

$$K(x, x') = \exp\left(-\frac{||x - x'||^2}{l^2}\right) \propto \frac{\text{Distance between points}}{\text{Lengthscale}}$$

GP mean prediction with small lengthscales

GP mean prediction with large lengthscale for input 2 is irrelevant)

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• GP hyperparameters are generally estimated by maximizing the marginal likelihood (Empirical Bayes)

$$\hat{\theta}_{MLE} = \arg \max - \frac{1}{2} \mathbf{y} [\mathbf{K} + \sigma_n^2 \mathbf{I}]^{-1} \mathbf{y}^T - \frac{1}{2} \log |\mathbf{K} + \sigma_n^2 \mathbf{I}| - \frac{N}{2} \log 2\pi$$
Data fit Complexity

- Being more Bayesian, GP hyperparameters can be treated as random variables with a prior distribution
- We can use Maximum A Posteriori (MAP) estimation as a way to encode more prior knowledge

$$\hat{\theta}_{MAP} = \arg \max - \frac{1}{2} \mathbf{y} [\mathbf{K} + \sigma_n^2 \mathbf{I}]^{-1} \mathbf{y}^T - \frac{1}{2} \log |\mathbf{K} + \sigma_n^2 \mathbf{I}| - \frac{N}{2} \log 2\pi + \log p(\theta)$$
Prior hyper-parameter
Data fit
Complexity
Prior hyper-parameter
probability

Being fully Bayesian, we sample from the hyperparameter prior using Markov Chain Monte Carlo (MCMC)

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Given the large number of factors, we can make two prior assumptions about the system under study [2,3] ٠



Inference: MAP estimation

Inference: MCMC

[2] Hvarfner et al, 2024, Vanilla Bayesian Optimization Performs Great in High Dimensions, ICML [3] Eriksson, Jankowiak, 2021, High-Dimensional Bayesian Optimization with Sparse Axis-Aligned Subspaces, UAI **General Business**

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- Acquisition functions quantify the value of information (regarding optimum), conditioned on prior beliefs and data
- Exploration (gather new information) is combined with exploitation (optimize given current knowledge)
- In this work, we rely on the log-Expected Improvement acquisition function, using Botorch Python package [4]



[4] Balandat et al 2020, BoTorch: A Framework for Efficient Monte-Carlo Bayesian Optimization, NEURIPS

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Results

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- D-optimal screening design can lead to different categorical level combinations with identical D-optimality
- To assess variability, 5 different repetions of sequential DOE are obtained by generating different D-optimal designs
- Different designs lead to different conclusions about factor importance
- I optimal design requires 39 to 44 additional runs for a total of 83 to 92 runs



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- 20 different BO repetions with 48 experiments, each with 10 initial experiments selected using random sampling
- All algorithms quickly converge near to the global optimum
- Best models: GP-SAAS for OHE, GP-DSP for DFT encodings



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- BO algorithms lead to better observed yields than both D and I optimal designs
- None of the 5 DOE repetions find the global optimum of 100% yield
- BO models lead to better results than final DOE models, while using only 48 runs (same as D-optimal screening design)



- Full dataset (1728 samples) is used to assess global prediction quality in test scenarios
- GPs use all categorical factors, LASSO model only uses a subset of factors
- GP-DSP-DFT leads to slighly better overall predictions and better predictions around the optimum

Metric for global prediction quality (1728 samples)	LASSO (44 training samples)	GP-DSP-DFT (48 training samples)	GP-SAAS-OHE (48 training samples)
R squared	0,32	0,34	0,36
Root mean squared error	20,2	19,5	16,0



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Conclusions

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5. Conclusions



- BO leads to faster convergence than sequential DOE on the same modeling basis (OHE)
- DFT encodings can improve optimization performance **291 covariates are efficiently used within BO**
- BO is an efficient appproach to experimental design with many categorical factors:
 - Faster convergence than sequential DOE
 - Avoid variability and uncertainty in the assessing important factors
 - Easier human implementation by avoiding screening stage and restrictive modeling assumptions
- We are currently exploring ways to include DOE principles that could improve BO even further

Thank you for your attention

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