Model-based Design for GPs

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- Model-free designs choose space-filling X_n. Without knowing much about the response surface you intend to model a priori, a space-filling design (i.e. LHS, maximin, minimax) represents a good choice.
- Recall if the response surface is linear, observations at boundaries are optimal, because they maximize leverage, minimize s.e. of $\hat{\beta}$.
- For a Gaussian Process (GP) response surface, we can choose samples optimal in some statistical sense.

"The best time to plan an experiment is after you've done it." - R. A. Fisher

• Sequential design helps avoid over-leveraging of prior beliefs before data collection.

- One-batch approach
 - Maximum entropy design (Maxent)
 - Minimum predictive uncertainty (IMSPE)
- Sequential approach
 - Active learning MacKay (ALM)
 - Active learning Cohn (ALC)

• Model assumption:

$$\begin{aligned} Y &= f(X) + \epsilon \\ \epsilon_i &\sim N(0, \tau^2 g) \\ f &\sim GP(0, \tau^2 K), K^{ij} = C_{\theta}(x_i, x_j) \\ \Rightarrow Y \mid X, g, \tau^2, \theta \sim N(0, \tau^2 (K + gI)) \end{aligned}$$

• Maximize the entropy of the marginal of Y w.r.t X_n :

$$-E\{\log p(Y|X_n,g,\tau^2,\theta)\}$$

 \Rightarrow Equivalent to maximizing $|K_n|$

• Most informative for Bayesian learning

Maxent: Algorithm

```
library(plgp)
maxent <- function(n, m, theta=0.1, g=0.01, T=100000)</pre>
  if(length(theta) == 1) theta <- rep(theta, m)</pre>
  X <- matrix(runif(n*m), ncol=m)</pre>
  K <- covar.sep(X, d=theta, g=g)</pre>
  ldetK <- determinant(K, logarithm=TRUE)$modulus</pre>
  for(t in 1:T) {
    row <- sample(1:n, 1)</pre>
    xold <- X[row,]</pre>
    X[row,] <- runif(m)
    Kprime <- covar.sep(X, d=theta, g=g)</pre>
    ldetKprime <- determinant(Kprime, logarithm=TRUE)$modulus</pre>
    if(ldetKprime > ldetK) { ldetK <- ldetKprime</pre>
    } else { X[row, ] <- xold }</pre>
  return(X)
```

Figure: Naive algorithm for maxent

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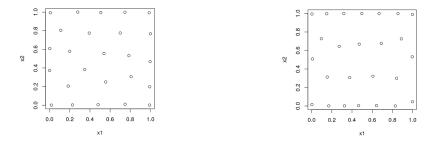


Figure: Maxent design under isotropic variance (left) and varying lengthscales (right)

- Strengths: adjust spread of different dimensions, theoretical guarantee.
- Limitations: design points cluttered at boundaries, few unique settings

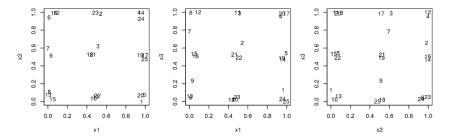


Figure: Projections of pairs of inputs involved in a 3d maximum entropy design

- Strengths: adjust spread of different dimensions, theoretical guarantee.
- Limitations: design points cluttered at boundaries, few unique settings, projections into lower dimensions don't have uniformity.

• Update $\log |K_{n+1}|$

$$\log |K_{n+1}| = \log |K_n| + \log v_n(x_{n+1})$$

where $v_n(x_{n+1}) = 1 + g - k_n(x_{n+1})^T K_n^{-1} k_n(x_{n+1})$
$$= \frac{\sigma_n^2(x_{n+1})}{\hat{\tau}_n^2}$$
$$\Rightarrow O(n^2)$$

- Update scale-free predictive variance $v_{n+1}(x)$
- Update precision matrix

Space-filling: spread points over the input space of interest (not related to prediction)

IMSPE: enhance prediction accuracy, interested in a sub-region of input space (local IMSPE, weighted IMSPE)

Model assumption:

$$Y(x) = \sum_{i=1}^{p} f_i(x)\beta_i + Z(x) = f^{T}(x)\beta + Z(x),$$
(1)

where Z(x) is a GP with Gaussian correlation function R(.).

• Minimize mean-sqared prediction error (MSPE):

$$MSPE\left(\mathbf{x}_{0}, \mathbf{X} \mid \sigma_{Z}^{2}, \boldsymbol{\rho}\right) = E_{Y}\left\{\left(Y(\mathbf{x}_{0}) - \widehat{\mathbf{y}}(\mathbf{x}_{0})\right)^{2}\right\}$$
$$= \sigma_{Z}^{2}\left(1 - \left[\boldsymbol{f}_{0}^{T} \ \boldsymbol{r}_{0}^{T}\right] \begin{bmatrix} \mathbf{0} \ \boldsymbol{F}^{T} \\ \boldsymbol{F} \ \boldsymbol{R} \end{bmatrix}^{-1} \begin{bmatrix} \boldsymbol{f}_{0} \\ \boldsymbol{r}_{0} \end{bmatrix}\right),$$

where F is known regressor, R_{ρ} is Gaussian correlation matrix, y^n is training outputs.

• generalized A-optimality: min trace of inverse of info matrix

• (known ρ)A local IMSPE: integrate out x_0

$$IMSPE(X|\sigma_Z^2,\rho) = \int_{[0,1]^d} MSPE(x_0, X|\sigma_Z^2,\rho) dx_0$$
(2)

Specially, if GP has constant mean (i.e. F are 1's), depends on ρ only

min
$$IMSPE(\ldots | \sigma_Z^2, \rho) = min IMSPE(\ldots | 1, \rho)$$

• (unknown ρ) A weighted IMSPE (W-IMSPE):

$$W(X|\pi) = \int_{[0,1]^d} IMSPE(X|1,\rho)\pi(\rho)d\rho$$
(3)

IMSPE: Computation

- Closed form: IMSPE with a known ρ ; rectangular input space and certain covariance kernels.
- Numeric approximation: W-IMSPE no available closed form
- Approximation: quasi Monte Carlo numerical integration based on a low discrepancy sequence

$$W(X|\pi) = \int_{[0,1]^d} IMSPE(X|1,\rho)\pi(\rho)d\rho \tag{4}$$

• (Leatherman et. al., 2018)

$$W_{a}(X|\pi) = \frac{1}{2^{k}} \sum_{j=1}^{2^{k}} IMSPE(X|1,\rho_{j})\pi(\rho_{j}),$$
(5)

where ρ_j is 2^k -point Sobol sequence. Modification: 1) adaptive k; 2) PSO algorithm to choose starting point. (Gramacy, retrieved 2021)

- other reference grids such as poor-man's quadrature or random reference grid.
- cons: not off-boundary; discrete or mixed continuous-discrete optimization
- using random reference grid for non-regular space
- improvements: a larger reference set, more stochastic exchange proposals, sequential design adaption, etc.

IMSPE: Comparison with space-filling design

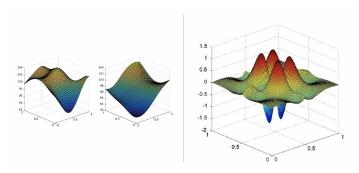


Figure: Example surfaces

- Smooth "stationary" surfaces, IMSPE-based methods are recommended
- Functions with pronounced non-stationary activity near the "middle" of the input domain: space-filling LHDs and MaxPro are recommended

- Similar to maxent or maximin
- avoid boundary of input space (sites at boundary don't cover space efficiently)
- In higher input dimension, more "off the boundary"

Sequential design/ Active learning

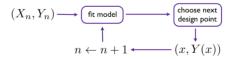


Figure: Diagram of Sequential design/active learning

- Assume a flexible surrogate, e.g., a GP model with unknown. hyperparamters
- Require outputs y ~ f(x), a choice of initial design size n and final size N, and criterion J(x) to choose to next point.

Then

- **()** Fit the surrogate (hyperparameters) using $D_n = (X_n, Y_n)$, e.g., via MLE.
- 2 Choose $X_{n+1} = \operatorname{argmax}_{x \in \mathcal{X}} J(x) | D_n$.
- Observe the response by running a new simulation to get y_{n+1}, and update D_{n+1}.

- More practical: In many situations, selecting one design point at a time works better than static, single-batch design
 - Single-batch design is sensitive to hyperparameters, while in sequential design, could **update hyperparameters** for each run.
 - ② Data measurements are relatively expensive or slow, and we want to know where to look next so as to learn as much as possible.
 - O There is an immense amount of data and we wish to select a subset of data points that is most useful for our purposes.
- The sample size N need not to be fixed, and information gain for each new data point is available.
 - **0** Omit the data points that are expected to be least informative
 - Form a stopping rule, so that we could decide whether to gather more data, given a desired exchange rate of information gain per measurement (Lindley 1956).

Setup:

Start with a LHS in 2d of size $n_0 = 12$ with $f(x_1, x_2) = x_1 \cdot e^{-x_1^2 - x_2^2} + N(0, 0.01^2).$

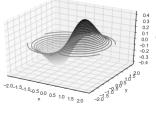


Figure: Function f

- ② Create a testing grid and saves true (noiseless) responses at those locations.
- Galculate RMSE to see out-of-sample progress over iterations of design acquisition

Criterion J(x) is predictive variance $\sigma_n^2(x)$ in ALM.

- Predictive variance produces sausage-shaped error-bars, so it must have many local maxima.
- The number of local maxima could grow linearly in sample size n. Optimizing globally over that surface presents challenges
- Solution Use the library-based local solver in R, "optim" with method="L-BFGS-B".

Thus, we adopt the multi-start scheme.

Multi-start Scheme

We "design" a collection of starting locations placed in parts of the input space known to have high variance.

```
xnp1.search <- function(X, gpi, obj=obj.alm, ...)</pre>
  start <- mvmaximin(nrow(X), 2, T=100*nrow(X), Xorig=X)</pre>
  xnew <- matrix(NA, nrow=nrow(start), ncol=ncol(X) + 1)</pre>
  for(i in 1:nrow(start)) {
    out <- optim(start[i,], obj, method="L-BFGS-B", lower=0,</pre>
      upper=1, gpi=gpi, ...)
    xnew[i,] <- c(out$par, -out$value)</pre>
  solns <- data.frame(cbind(start, xnew))</pre>
  names(solns) <- c("s1", "s2", "x1", "x2", "val")</pre>
  return(solns)
```

Figure: Function for searching x_{n+1}

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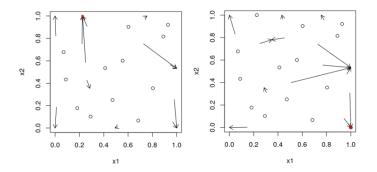


Figure: First/second iteration of ALM search. Each arrow represents an origin and outcome of multi-start exploration of predictive variance. Variance-maximizing location is indicated as a red dot.

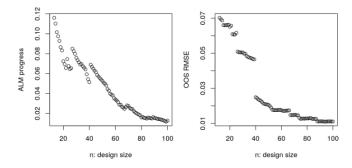


Figure: Maximum variance (left, lower is better) and out-of-sample RMSE (right) over 100 ALM acquisitions.

Progress metrics and RMSE are starting to level off in the later 35 iterations or so.

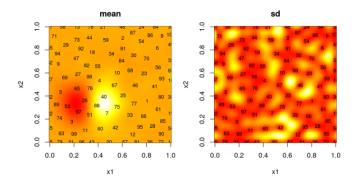


Figure: Predictive mean (left) and standard deviation (right) after ALM-based sequential design.

Observe dense coverage along the boundary since variance is high along the boundary.

- ALM can misbehave, especially if the starting design is unlucky to miss strong signal in the data
- 2 ALM doesn't recognize that acquisitions impact predictive equations globally.
- Potentially ignoring a fatter regions where uncertainty may cumulatively be much larger.
- Variance is high along the boundary because there are fewer data points nearby, so we end up with lots of points on the boundary.

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 - Global: integrate over the whole space;

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- Might be better to consider how much reduction in posterior variance can be obtained by adding an extra point.
- Question: where should the reduction be measured?
- Two extremes:
 - Global: integrate over the whole space;
 - Local: at specific reference point(s).

- Cohn (1994) suggests such an acquisition heuristic in a nonparametric regression context for neural networks.
- Seo et al. (2000) adapt Cohn's ideas to Gaussian Process and called it ALC.

- How does it work?
- Recall that predictive variance follows

$$\sigma_n^2 = \hat{\tau}_n^2 [1 + \hat{g}_n - k_n^{\mathrm{T}}(x) K_n^{-1} k_n(x)], \text{ where } k_n(x) \equiv C_{\hat{\theta}_n}(X_n, x).$$

• The deduced variance

$$\tilde{\sigma}_{n+1}^2 = \hat{\tau}_n^2 [1 + \hat{g}_n - k_{n+1}^{\mathrm{T}}(x) \mathcal{K}_{n+1}^{-1} k_{n+1}(x)], \text{ where } k_{n+1}(x) \equiv C_{\hat{\theta}_n}(X_{n+1}, x).$$

 The ALC criterion is the average reduction in variance from n → n + 1 measured through a choice of x_{n+1}:

$$\Delta \sigma_n^2(x_{n+1}) = \int_{\mathcal{X}} [\sigma_n^2(x) - \tilde{\sigma}_{n+1}^2(x)] \, \mathrm{d}x.$$

• The criterion must be solved in each iteration of sequential design.

$$x_{n+1} = \arg\min \tilde{\sigma}_{n+1}^2(x), x \in \mathcal{X}.$$

- Closed form when \mathcal{X} is rectangular.
- Often in practice approximated by a reference set.

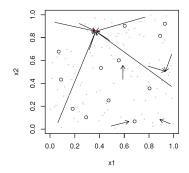


Figure: First iteration of ALC search. Gray dots denote reference locations.

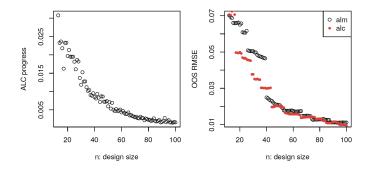
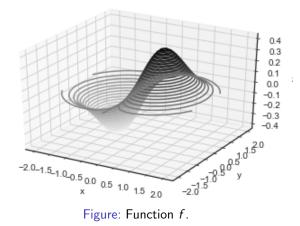


Figure: Progress in ALC sequential design in terms of integrated reduction in variance (left, lower is better) and out-of-sample RMSE (right), with comparison to ALM.

Recall what the true function looks like.



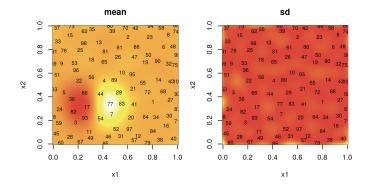


Figure: Predictive mean (left) and standard deviation (right) after ALC-based sequential design.

- Thinking about the hyperparameters what point can we select in order to estimate the hyperparameters more accurately?
- Criterion: maximize the Fisher Information.
- Does not lead to designs with the most accurate predictors.
- Hybrid approach:
 - FI learn hyperparameters
 - ALC prediction

Other Sequential Criteria - Fisher Information

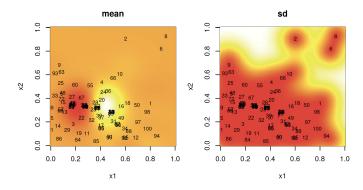


Figure: Predictive mean (left) and standard deviation (right) after FI-based sequential design.

Other Sequential Criteria - Fisher Information

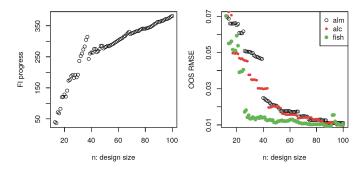


Figure: Progress in terms of FI (left, higher is better) and out-of-sample RMSE as compared to previous heuristics.