

Comment

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We have been running computer experiments related to semiconductor process design and recently switched over to the paradigm described by the authors. We have found it to be more flexible than response surface methodology in handling deterministic responses.

The Bayesian approach suggests how to interpolate, extrapolate, assess uncertainty and construct designs. To what extent do the Bayesian answers make sense, if one does not hold the prior belief? The authors cite several works in which connections are drawn between well accepted interpolation methods and various priors and give an example in which the uncertainty assessment is accurate. It would be very interesting if the uncertainty assessments were reasonably accurate for a large class of underlying functions. Have the authors investigated this point? We doubt that the Bayesian method will help in extrapolation (which we suspect should be avoided) and thus are worried that the optimal designs sometimes concentrate near the center of the design space.

Our main comments are directed at the design problem and at estimation of the parameters of the covariance model. Our applications have 5 to 10 input variables and a like number of outputs. The programs we use are fast enough to make it feasible to consider 50 or more runs.

Before addressing the design and estimation issues, we wish to point out that ideas from exploratory data analysis have a role to play in computer experimentation. The authors (with their coworkers) have plotted contours, trajectories and the additive main effects (mentioned in Section 6) of the response functions. We think their contributions are noteworthy and look forward to further developments. When there are many response variables, care should be taken in optimizing a functional of the responses without first considering the tradeoffs among competing goals. The approach taken in Sharifzadeh, Koehler, Owen and Shott (1989) is to evaluate the model functions at thousands of input points and to explore the resulting

data set with interactive graphics, in this case S (Becker, Chambers and Wilks, 1988).

DESIGN ISSUES

In the authors' Figure 1, the design points are all quite close to the center. We share the misgivings of the authors, suspecting that this leads to a robustness problem. Extrapolation by conditional expectation depends to a far greater degree on the covariance function used than does interpolation. Thus outside of the convex hull of the data, the predicted values will depend strongly on hard-to-verify properties of the model.

We have been using low discrepancy sequences, mentioned in Section 7.5, as designs. These designs are constructed so that the empirical measure of the design points is close in a Kolmogorov-Smirnov sense to the uniform measure on the cube. These should be good designs in the case of large θ , when estimation is difficult. Johnson, Moore and Ylvisaker (1988) characterize the optimal designs in the large θ limit. Minimizing the maximum distance from a point in the cube to a design point leads to their version of G optimality and maximizing the minimum distance between two sample points leads to their version of D optimality. Low discrepancy sequences (such as Halton-Hammersley sequences) tend to have small, but not minimal, maximum distances from points in the cube.

We have found that sometimes some of the θ_j appear quite small while others are large. That is a response variable is heavily dependent on a few of the d inputs and not very sensitive at all to the others. We may not know in advance which input variables are the important ones or, more commonly, each output variable may depend most strongly on a different small set of inputs. This opens up the possibility of reducing the dimension of the problem by considering the response as a function of the most important inputs, possibly with some noise due to the other inputs. For instance, in our first experiment the thickness of a layer of SiO₂ only depended on the oxidation temperature. Unfortunately, our design (an all-bias design) only used three distinct values of the temperature in 43 runs. If our experiment had had 43 nearly equispaced temperature values, the results would have been more informative.

Low discrepancy designs have the added benefit that when projected onto a cube defined by a subset of the original variables they are still nearly uniform.

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Thus if the dimension can be reduced, the design in the remaining dimensions is still reasonably good. The optimal designs depicted in Johnson, Moore and Ylvisaker (1988) do not tend to project uniformly.

We prefer the sequences of Faure (1982) to the Halton-Hammersley sequences. The Halton-Hammersley sequences are usually based on the first d prime numbers, whereas Faure uses the same prime number (the smallest prime $r \geq d$) on each axis. When $n = r^k$, the Faure sequences exercise each input variable in much the same way Latin hypercube designs do. Moreover for $k \geq 2$ they exercise pairs of input variables in that, for any given pair of inputs, one can partition their domain into r^2 squares and find r^{k-2} points in each square. Similarly there are equidistribution properties for three or more axes. The equidistribution properties of the Halton-Hammersley sequences are different for each marginal subcube, depending on the associated primes. We have found that with $n = r^2$ and $r = 5$ or 7 that the Faure sequences appear to lie on planes in three dimensions. This is alleviated by replacing each digit b in the base r representation of the Faure sequence by $\sigma(b)$ where σ is a permutation of $0, \dots, r - 1$. The permutation does not alter the equidistribution properties. One can inspect three-dimensional scatterplots to make sure that a given permutation is effective.

PARAMETER ESTIMATION

We would like to mention a quick way of estimating $\theta_1, \dots, \theta_d$ in the covariance given by the authors'

equation (9) with $p = 1$. When the function $Y(x)$ is nearly additive, we can estimate the main effects using scatterplot smoothers. This corresponds to the inner loop of the ACE algorithm in Breiman and Friedman (1985). Let g_j denote the estimate of the j th main effect. A very smooth $g_j(\cdot)$ is evidence that θ_j is small and a rough $g_j(\cdot)$ suggests that θ_j is large. The roughness may be assessed by $\mathcal{R}_j = \sum_{i=1}^m (g_j(i/m) - g_j((i-1)/m))^2$ where the domain of g_j has been rescaled to $[0, 1]$. The expected value of \mathcal{R}_j may be expressed in terms of θ_1 through θ_d , for fixed σ . The d equations in d unknowns can be solved iteratively. The likelihood can be used to choose between the answers from several different values of m . This avoids a high dimensional search for $\theta_1, \dots, \theta_d$. The first time we tried it, we got better parameter values (as measured by likelihood) than we had found by searching. Alternatively it suggests starting values for such a search.

ADDITIONAL REFERENCES

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Comment

Anthony O'Hagan

The authors are to be congratulated on their lucid and wide-ranging review. Like others before, I have independently rediscovered many of the ideas and results presented here. I therefore sincerely hope that the greater prominence given to those ideas and results by this excellent paper will enable future researchers to start well beyond square one. I first have some comments concerning the derivation of the basic estimator (7), and I will then discuss the model and the practical implementation of the methods from my own experience.

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The authors mention three derivations of (7). In a classical framework, it is the MLE if the process $Z(\cdot)$ is Gaussian, and relaxing this assumption it is the BLUP, minimizing (2). Thirdly, it is the posterior mean of $Y(x)$ in a Bayesian analysis with a Gaussian $Z(\cdot)$ and a uniform prior on β . It is first worth pointing out that with a proper multivariate normal prior $\beta \sim N(b, B)$ and known σ^2 the posterior mean of $Y(x)$ has the same form as (7), but with $\hat{\beta}$ replaced by the posterior mean of β , i.e.,

$$\tilde{\beta} = (F'R^{-1}F + \sigma^2B^{-1})(F'R^{-1}F\hat{\beta} + \sigma^2B^{-1}b).$$

The interpretation of (7), as comprising the fitted regression model plus smoothed residuals, still holds.

We can also dispense with normality in the Bayesian framework, using a similar device to (2). The