# Sequential, robust design strategies

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# Doug Wiens University of Alberta

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### Approximate regression models

• Experimenter fits a response  $\hat{Y}(\mathbf{x}) = f\left(\mathbf{x}; \hat{\boldsymbol{\theta}}\right)$  by regression, when in fact

$$E[Y|\mathbf{x}] pprox f(\mathbf{x}; oldsymbol{ heta})$$
 .

- The points x<sub>i</sub> at which Y will be observed are to be chosen with an eye to protection against a misspecified response function.
- Best fitting parameter is

$$oldsymbol{ heta}_0 = rgmin \int_S \left\{ E([Y|\mathbf{x}] - f(\mathbf{x}; oldsymbol{ heta}) 
ight\}^2 d\mathbf{x}$$
  
for  $\mathbf{x} \in \mathcal{S}$  ("design space").

• Put  $g(\mathbf{x}) = E[Y|\mathbf{x}] - f(\mathbf{x}; \theta_0)$ ; then (additive errors)

$$Y(\mathbf{x}) = f(\mathbf{x}; \boldsymbol{\theta}_0) + g(\mathbf{x}) + \varepsilon.$$

PROBLEM: Choose a design  $\xi$  (= a measure placing mass  $n^{-1}$  at selected points  $\mathbf{x}_1, ..., \mathbf{x}_n \in S$ ) so as to minimise loss due to:

- random variation; depends only on  $\xi$
- bias (of Ŷ(x) as estimate of E[Y|x]; depends on (g, ξ))

Loss: Integrated MSE of the predictions

$$\begin{aligned} \mathcal{L}(g,\xi) &= \int_{\mathcal{S}} E\left[\left\{\hat{Y}(\mathbf{x}) - E(Y|\mathbf{x})\right\}^{2}\right] d\mathbf{x} \\ &= \int_{\mathcal{S}} VAR\left[\hat{Y}(\mathbf{x})\right] d\mathbf{x} \\ &+ \int_{\mathcal{S}} \left\{E\left[f\left(\mathbf{x};\hat{\boldsymbol{\theta}}\right) - f\left(\mathbf{x};\boldsymbol{\theta}_{0}\right) - g(\mathbf{x})\right]\right\}^{2} d\mathbf{x} \end{aligned}$$

- Find ξ<sub>0</sub> = arg min L(g, ξ) after
  (i) maximising over g (= E [Y|x] f (x; θ<sub>0</sub>)); or
  (ii) estimating g.
- Sequential strategy may be called for, in either case
- $\hat{\theta}$  can be LSE, or M-estimate (with  $\sigma^2$  replaced by, e.g.,  $\sigma^2 E\left[\psi^2\right] / (E\left[\psi'\right])^2$ ).

NONLINEAR REGRESSION (with Sanjoy Sinha):

Fit  $E[Y|\mathbf{x}] = f(\mathbf{x}; \boldsymbol{\theta}_0)$  when in fact this is only approximate, e.g.

 $f(x; \theta_0) = \theta_0 e^{-\theta_1 x}$  but  $E[Y|\mathbf{x}] = \frac{\theta_0 x}{\theta_1 + x}$ .

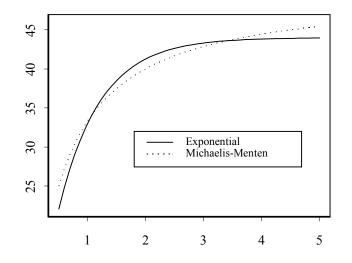


Figure 1: E[Y|x] is Michaelis-Menten with  $\theta = (50, .5)^T$ ; best-fitting exponential is  $f(x; \theta_0)$  with  $\theta_0 = (44, 1.39)^T$ .  $(\theta_0 = \arg \min \int_{.5}^{5} \{E([Y|\mathbf{x}] - f(\mathbf{x}; \theta)\}^2 d\mathbf{x}.)$ 

$$g(\mathbf{x}; \boldsymbol{\theta}_0) = E[Y|\mathbf{x}] - f(\mathbf{x}; \boldsymbol{\theta}_0)$$

Asymptotic MSE matrix is  $MSE_N(\theta_0) = M_N^{-1}(\theta_0) \left\{ Q_N(\theta_0) + b_N(\theta_0) b_N^T(\theta_0) \right\} M_N^{-1}(\theta_0),$ where  $\mathbf{z}(\mathbf{x}; \theta) = \partial f(\mathbf{x}; \theta) / \partial \theta$  and where

$$egin{aligned} \mathbf{M}_N(m{ heta}) &=& \sum\limits_{i=1}^N \mathbf{z}(\mathbf{x}_i;m{ heta}) \mathbf{z}^T(\mathbf{x}_i;m{ heta}), \ \mathbf{Q}_N(m{ heta}) &=& \sum\limits_{i=1}^N \mathbf{z}(\mathbf{x}_i;m{ heta}) \sigma^2(\mathbf{x}_i) \mathbf{z}^T(\mathbf{x}_i;m{ heta}), \ \mathbf{b}_N(m{ heta}) &=& \sum\limits_{i=1}^N \mathbf{z}(\mathbf{x}_i;m{ heta}) g(\mathbf{x}_i;m{ heta}). \end{aligned}$$

Loss is IMSE:

$$\begin{split} \mathcal{L}(g,\xi) &= \int_{\mathcal{S}} E\left[\left\{\hat{Y}(\mathbf{x}) - E(Y|\mathbf{x})\right\}^2\right] d\mathbf{x} \\ &\approx tr\left[\mathsf{MSE}_N(\theta_0) \cdot \mathbf{A}(\theta_0)\right] + \int_{\mathcal{S}} g^2(\mathbf{x};\theta_0) d\mathbf{x}, \\ \end{split}$$
where  $\mathbf{A}(\theta) = \int_{\mathcal{S}} \mathbf{z}(\mathbf{x};\theta) \mathbf{z}^T(\mathbf{x};\theta) d\mathbf{x}.$ 

Sequential approach. Given  $\{\mathbf{x}_i, Y_i\}_{i=1}^N$ : (i) Compute  $\hat{\boldsymbol{\theta}}_N$  and estimates of  $g(\mathbf{x})$ ,  $\sigma^2(\mathbf{x})$ . (ii) Using these estimates, estimate  $\Delta_{N+1}(\mathbf{x}) =$ increase in  $\mathcal{L}$  if the next design point is  $\mathbf{x}$ . (iii) Choose  $\mathbf{x}_{N+1} = \arg \min \Delta_{N+1}(\mathbf{x})$ .

Estimate  $g(\mathbf{x})$  by smoothing the residuals (cubic spline in 1-dimensional; generalised additive model for higher dimensions).

Asymptotic results hold for sequentially chosen design points - Sinha and Wiens (2002).

CLINICAL TRIALS: Subjects are assigned to one of p treatment groups. Covariates x are measured and treatment assignments made, according to a random mechanism.

Optimal assignment probabilities

$$\mathsf{Pr}\left(\mathsf{treatment}\ i|\mathbf{x}\right) = \rho_i(\mathbf{x})$$

are to be determined.

Post treatment response to treatment is

$$Y = \theta_i + \mathbf{z}^T(\mathbf{x})\boldsymbol{\phi} + g_i(\mathbf{x}) + \sigma_i\varepsilon$$

for regressors z(x), error variances  $\sigma_i$ , response errors  $g_i(x)$ .

Design 
$$\xi = \{\rho_1, ..., \rho_p\}.$$

Let  $W_{p-1 \times p}$  have rows which are mutually orthogonal and orthogonal to 1. We estimate a complete set  $W\theta$ of contrasts of the treatment effects  $\{\theta_i\}_{i=1}^p$ .

Loss is

$$\mathcal{L}\left(\rho_{1},...,\rho_{p}\right) = \lim_{n \to \infty} \left| nMSE\left(\mathbf{W}\hat{\boldsymbol{\theta}}\right) \right|.$$

 Heckman (1987) - similar approach; different neighbourhood structure. Under realistic conditions *constant* assignment probabilities were found to be optimal. It turns out that constant probabilities

 $\rho_i(\mathbf{x}) \equiv r_i$ 

minimize the COV part of MSE.

Optimal  $\{r_i\}_{i=1}^p$  are those which

minimise 
$$rac{\sum \left(r_i/\sigma_i^2\right)}{\prod \left(r_i/\sigma_i^2\right)},$$

subject to  $\{r_i\}_{i=1}^p$  being a probability distribution.

When p = 2,

$$r_i = \frac{\sigma_i}{\sigma_1 + \sigma_2}.$$

Sequential assignments. Adjust the (asymptotically) variance minimising  $\{r_i\}_{i=1}^p$ , while also minimising variance and bias in finite samples.

Suppose there are L levels of the (grouped) covariates  $\mathbf{x}^{(1)}, ..., \mathbf{x}^{(L)}$ . If n assignments have been made, and the  $(n+1)^{th}$  subject arrives with covariates  $\mathbf{x}_*$ , then assign to treatment k with probability

$$P(k|\mathbf{x}_*) \propto \hat{r}_k d_k^* b_k^*,$$

where:

(i)  $\hat{r}_k$  is the optimal r, with the  $\sigma_i$  estimated.

(ii)  $d_k^*$  measures the reduction in  $\left| COV \left( \mathbf{W} \hat{\boldsymbol{\theta}} \right) \right|$  resulting from an assignment to treatment k.

(iii)  $b_k^*$  is inversely proportional to the (finite sample) bias<sup>2</sup> of  $\hat{\theta}$ , resulting from an assignment to treatment k.

$$P\left(k|\mathbf{x}_{*}
ight) \propto \hat{r}_{k}d_{k}^{*}b_{k}^{*}$$

Similar to Atkinson (1982) who takes  $P(k|\mathbf{x}_*) \propto d_k^*$  (assuming no bias, and that all  $\sigma_i^2$  are equal).

Computation of  $b_k^*$  requires  $\hat{g}_1(\mathbf{x}), ..., \hat{g}_p(\mathbf{x})$ ; an *ad hoc* estimate is the adjusted residual

$$\hat{g}_i(\mathbf{x}^{(l)}) = sign\left(\tilde{e}_{i,l}\right) \left(\tilde{e}_{i,l}^2 + \frac{\hat{\sigma}_i^2}{n_{i,l}}\right)^{1/2},$$

where  $n_{i,l} = \#$  of assignments of  $\mathbf{x}^{(l)}$  to group *i*;  $\tilde{e}_{i,l} =$  median of corresponding residuals.

#### SPATIAL STUDIES

- Observe  $Y(\mathbf{t}) = X(\mathbf{t}) + \varepsilon(\mathbf{t})$  at locations  $\mathbf{t} \in \mathcal{T} \subset \mathbb{R}^d$ .
- $X(\mathbf{t})$  random:  $X(\mathbf{t}) = E[X(\mathbf{t})] + \delta(\mathbf{t})$ .
- $E[X(t)] \approx \mathbf{z}^T(t) \boldsymbol{\theta}$  for regressors  $\mathbf{z}(t)$
- VAR [ε(t)] = f(t) only approximately known (assumed constant?)
- $COV[\delta(t), \delta(t')] = g(t, t')$  only approximately known (assumed isotropic?)
- Choose n locations from T (with N sites) so as to minimise the MSE of the predictions, maximised over neighbourhoods of the assumed f, g and regression model.

#### NEXT:

- Sequential choice of sites?
- Simulated annealing?

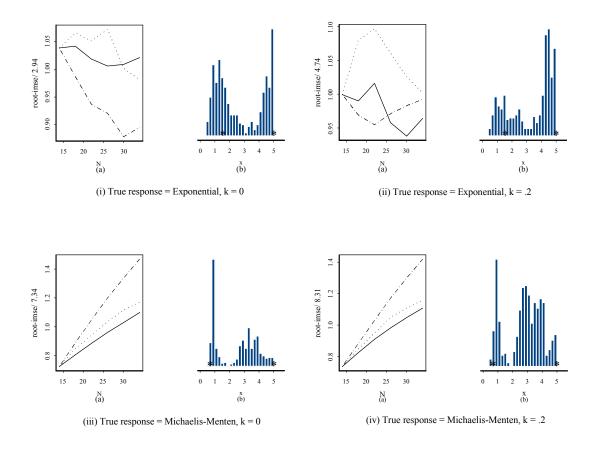


Figure 2: Fitted response is exponential, true response is either exponential or Michaelis-Menten;  $n_0 = 10$  equally spaced sites chosen initially, with  $r_0 = 3$  replicates at each. Then  $n_1 = 6$  additional sites chosen sequentially, with  $r_1 = 4$  replicates at each. (a) Average (over 100 sample paths) values of  $(N \cdot IMSE)^{1/2}$  for sequential (----), uniform (...) and D-optimal (-..-) designs. Variance function is  $\sigma^2(x) = 1 + .2(x - .5)^2$ . (b) Probability histogram of all points chosen by the 100 sequential designs; asterisks are at the average sites of the D-optimal designs.

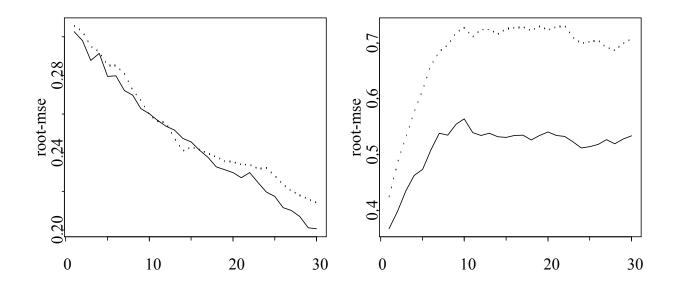


Figure 3: Root-mse of estimated treatment effects versus new subjects; average of 200 simulated runs. Two treatments, two covariates  $X_1, X_2$ . Heteroscedastic errors:  $\sigma_1^2 = 1$ ,  $\sigma_2^2 = 1/4$ . Dotted line is Atkinson's method modified for heteroscedasticity:  $P(k|\mathbf{x}_*) \propto \hat{r}_k d_k^*$ ; solid line is the robust method. Left:  $g_1(\mathbf{x}) = g_2(\mathbf{x}) \equiv 0$  (fitted model correct). Right:  $g_i(\mathbf{x}) \propto (-1)^i x_1 x_2$ .

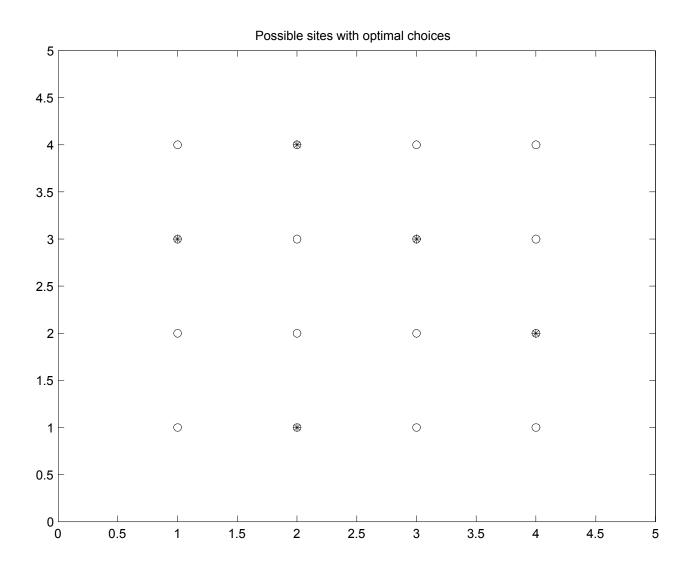


Figure 4:  $4 \times 4$  grid of possible locations; 5 sites chosen to minimise trace of MSE matrix. Fitted model exact: constant measurement errors, isotropic covariance function exp $(-.2 ||\mathbf{t} - \mathbf{t'}||)$ , regressors  $\mathbf{z}(\mathbf{t}) = (1, t_1, t_2)^T$ .

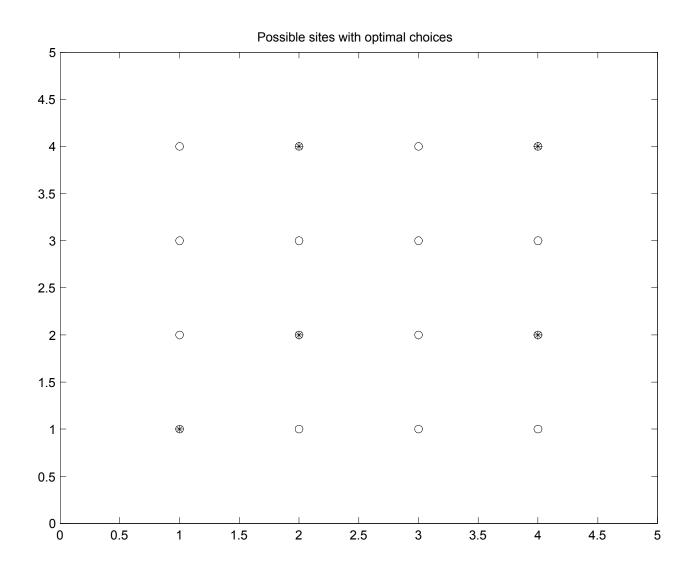


Figure 5: Same fitted model, but loss is maximised over neighbourhoods of the model, then minimised over choices of locations.