Optimal Designs for Estimating Critical Effective Dose Under Model Uncertainty in a Dose Response Study

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Abstract

Toxicologists have been increasingly using a class of models to describe a continuous response in the last few years. This class consists of nested nonlinear models and is used for estimating various parameters in the models or some meaningful function of the model parameters. Our work here is the first to address design issues for this popular class of models among toxicologists. Specifically we construct a variety of optimal designs under model uncertainty and study their properties for estimating the critical effective dose (CED), which is model dependent. Two types of optimal designs are proposed: one type maximizes the minimum of efficiencies for estimating the CED regardless which member in the class of models is the appropriate model, and (ii) dual-objectives optimal design that simultaneously selects the most appropriate model and provide the best estimates for CED at the same time. We compare relative efficiencies of these optimal designs and other commonly used designs for estimating CED. To facilitate use of these designs, we have constructed a website that practitioners can generate tailor-made designs for their settings.

Keyword and phrases: compound optimal design, critical effect size, local optimal design, maximin optimal design, model discrimination, robust design.

1   Introduction

This paper addresses design issues for dose response studies in toxicology when the main outcome is continuous and it is not known a priori which model is an appropriate one to use. Under this situation, one may consider a class of plausible models within which we believe lies an adequate model for fitting the data at hand. The issues of interest are how to design to select the 'best' model from the class and at the same time to estimate the critical effective dose (CED) efficiently. The estimated CED is the dose that toxicologists use to estimate the dose that will result in a user-specified change in the continuous outcome after accounting for background noise. The user-specified change in the continuous outcome is usually expressed in terms of the critical effective size (CES), which is somewhat analogous to specifying 'alpha' in hypothesis testing in statistical inference.

Ideally, we want the design to be able to identify the correct model from the postulated class of models and also provide an efficient estimate for the CED, which is a function of the parameters in the identified model. In this paper, it is further assumed for simplicity that there is only one independent variable, the dose level. The design space is the range of dose levels of interest where the researcher selects the dose levels to observe the outcome. Throughout, we assume all design issues have to be decided in advance of the study and so sequential designs are not considered.

By design, the researcher has to select the number of dose levels from the design space to observe the continuous outcome, decide where these dose levels are and the number of replicates at each of these dose levels. Here the design space is the range of dose levels that the researcher wants to include in the study. We further generalize the design problem to one for finding continuous designs, meaning that we now view designs as probability measures on the design space. Continuous designs were proposed by

Kiefer in the late 1950s and have been shown to be much more amendable to analytical description and study than exact design. In our setup, we assume the total number of observations N for the whole study is pre-determined by cost or time and continuous designs are implemented by naturally rounding the possibly non-integer number of observations required at each dose to an integer number, subject to the number of observations at each dose sum to N.

Here is a simple illustration of a continuous design on the design space $[0, 25]$. Let ξ be a continuous design that takes half the observations at dose level 5 and half at dose level 20. We denote this design by writing $\xi = \{5, 20; 1/2, 1/2\}$; the first part denotes the two dose levels and the latter part denotes the corresponding proportion of observations to be taken at each of the dose levels. In the terminology of optimal design literature, the dose levels of the design ξ are called support points and the corresponding proportions are called weights. If $N = 20$, this implies the continuous design ξ takes 10 observations at dose level 5 and 10 observations at dose level 20. If $N = 25$, the same continuous design ξ can either take 12 observations at dose level 5 and 13 observations at dose level 20, or alternatively, 13 observations at dose level 5 and 12 observations at dose level 20. An optimal (continuous) design is one that maximizes or minimizes a given optimality criterion over all designs on the design space. Further details and motivations for working with continuous designs are given in the voluminous collection of papers by Kiefer and edited by Brown, et al. (1985). Optimal rounding procedures to convert a continuous design to an exact design for implementation are given in Rieder and Pukelsheim (1992).

Addressing design issues invariably requires model assumptions that specify how the mean outcome relates to the independent variable. Usually a specific functional form is assumed either from experts' opinions or from the science of the problem, see Gaylor

and Chen (1993), Catalano et al. (1993), Slob and Pieters (1998), Oscar (2004), Moerbeek, Piersma and Slob (2004), among many others. When it is problematic to specify a single model to describe the functional relationship between the mean outcome and the dose level, a common strategy is to work with a class of plausible nested models assumed to include the 'true' model. This class of models is usually arrived at after consultation with experts in the area. As an illustration, consider a simple class of nested models widely used in the study of enzyme kinetics. The class consists of just two models: the well known Michaelis-Menten and the Emax-model. The former is frequently employed in biochemistry and is described by

$$E(Y) = \frac{ax}{b+x}.$$

Here Y is the velocity of the enzyme kinetics and x is the concentration of the substract. The two parameters are a and b with the latter often referred to as the Michaelis-Menten constant. A more flexible model to study enzyme kinetics is the Emax model defined by

$$E(Y) = \frac{ax^h}{b+x^h}.$$

The extra parameter h in the Emax-model permits the shape of the response curve to be skewed and takes on different steepness as the concentration of the substract is varied. The challenges here from the design perspective are that we do not know at the onset which one of these two models is a more suitable model to use and it is well known that an optimal design can be very sensitive to model assumptions. If one considers the Michaelis-Menten alone, the optimal design for estimating the parameters a and b in this model does not allow one to estimate the parameter h in the Emax model and, if one assumes the Emax model holds, the optimal design for estimating all the three parameters h, a and b may be inefficient for only estimating a and b should the Michaelis-Menten model prove to be a more appropriate model. Such design problems are important and arise frequently in practice across disciplines. To our knowledge,

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only a couple of papers have tried to address such design questions seriously. A main reason for lack of research in this area is that there are serious technical difficulties involved, especially for nonlinear models.

A general strategy to address such design issues is to work with experts in the area and first identify a class of plausible modes that will most likely include the true model. The plausible models within this class should be nested allowing one to be built upon another; typically this means the 'largest' model has the largest number of parameters and the next 'largest' model is obtained from the 'largest' one by specifying one or more parameters equal to some user-selected fixed values. In our above illustration, our class of plausible models consists of the Emax and Michaelis-Menten models and it is clear that when $h = 1$, the Emax-model reduces to the Michaelis-Menten model. Once this class is identified, one works assuming the largest model holds. In our illustration, one seeks an efficient design to estimate h in the Emax model as accurately as possible and at the same time also have efficient estimates for the parameters a and b when the Michaelis-Menten model holds. Dette, Melas and Wong (2005) addressed this particular design problem and provided details.

The motivation for this work comes from repeated proposals recently in the toxicology literature to use a class of models to study a continuous outcome in toxicological studies [Moerbeek, Piersma and Slob (2004), Piersma et al. (2002), Woutersen et al. (2001), Slob (2002)]. In all these papers, the interest was only in estimation problems and so they did not consider design issues. As is typical in such publications, the rationale for the design employed in the study is not explained. Here are a few examples of designs used in toxicological studies and their outcomes. In Piersma et al. (2002), rats were prenatally exposed to diethylstilbestrol and the design had 16 animals in each of the 10 dose groups at 0, 1.0, 1.7, 2.8, 4.7, 7.8, 13, 22, 36 and 60 mg/kg body weight per

day. In Woutersen et al. (2001), rats were exposed to Rhodorsil Silane in a 28-day toxicity study and 3 designs were employed: the first one had 10 rats in each of the 7 dose groups (7x10 design) and the second had 5 rats in each of the 7 dose groups; the 7 dose groups were 0, 50, 150, 300, 450, 600 and 750 mg/kg body weight/day. The third had 10 rats in each of the 4 dose groups at 0, 50, 150 and 450 mg Rhodorsil Silane/kg body weight/day. There were many continuous outcomes in each of these studies. In Piersma et al. (2002), they included maternal body weight on gestation day 21, maternal serum estradiol concentration at gestation day 21, weights of fetuses at gestation day 21, immunological responses such as IgG and IgM to sheep red blood cell challenge and pup weights at days 1 and 21. In Woutersen et al. (2001), their main goals were to estimate various critical effective doses; these are doses that will result in a user-specified level of toxicity found in rats over the background noise. Outcomes of main interest were responses in haematology and clinical chemistry. In the discussion section, we comment on the performance of these designs.

In this paper, we develop optimal designs for identifying an appropriate model within the class of models and also at the same time provide reliable estimate for the critical effect dose (CED) using the selected model. The CED is a popular measure among toxicologists to estimate the dose level that will result in a user-specified anticipated change in the continuous outcome beyond the background noise. Design issues are always difficult to address and we begin first by considering local optimal designs because they are the easiest to construct for nonlinear models (Chernoff, 1953). These designs require the user to supply nominal values of the model parameters before the optimal design can be constructed. Nominal values represent the best guess for the true values of the set of parameters and are usually obtained either from prior similar experiments or experts' opinion. When model assumptions are mis-specified, it is well known that the resulting local optimal design can lose substantial efficiency. To overcome this

risk, we propose maximin optimal designs that have been shown to be robust to mis-specification of model assumptions in other settings , see for example, Biedermann, Dette and Pepelyshev (2006) and Dette et al. (2008). These maximin optimal design maximizes the minimum efficiency regardless which model in the class of models is the appropriate model. As such, these optimal designs provide some protection against picking a wrong model from the postulated class. In addition, we construct compound optimal design to account for the dual objectives of discriminating models and at the same time want the design to deliver a user-specified level of efficiency for estimating the CED.

In section 2, we describe the class of nonlinear models and the design criterion for estimating CED. We describe relationships among models in the class and provide local optimal design for estimating the CED for each member in the class. We also show how an optimal design constructed for a specific setup can be used to deduce the optimal design under another setup where assumptions on the design space and model parameters are different. In section 3, we construct maximin optimal designs and compound optimal designs for toxicology studies and assess their robustness properties to model mis-specification and their effectiveness for discriminating between models and estimating CED at the same time. We also compare performance of selected uniform designs that are intuitively appealing to practitioners. These designs take equal number of observations over a set of equally spaced dose levels. In section 4, we discuss four practical issues. First, we construct and compare corresponding optimal designs for log-normally distributed outcomes, which is another popular assumption used by toxicologists. Secondly, we investigate efficiencies of several designs used by toxicologists. Thirdly, we perform a simulation study to assess how our optimal designs perform in a real example with a relatively small sample size. Finally, we introduce the reader to our design web site that the reader can use freely to generate a variety

of optimal designs for a broad range of models frequently used in the biological sciences.

2 Model Uncertainty and CED-Optimality

Moerbeek, Piersma and Slob (2004), Woutersen et al. (2001), Piersma, et al. (2002) and Slob (2002) proposed and used the following class of models in several toxicological studies. The authors showed with justifications that the class of models is sufficiently flexible to accommodate typical continuous outcomes of interest in toxicological studies. For each of these models defined on the given design space $[0, T]$, Y is the response and t is the dose level; all parameters in the mean response are components of the $px1$-vector parameter θ .

$$(2.3) \qquad\qquad E(Y) = ae^{-bt} \text{ with } a > 0, \ b > 0,$$

$$(2.4) \qquad\qquad E(Y) = ae^{-bt^d} \text{ with } a > 0, \ b > 0, \ d \geq 1,$$

$$(2.5) \qquad\qquad E(Y) = a(c - (c-1)e^{-bt}) \text{ with } a > 0, \ b > 0, \ c \geq 0,$$

$$(2.6) \qquad\qquad E(Y) = a(c - (c-1)e^{-bt^d}) \text{ with } a > 0, \ b > 0, \ c \geq 0, \ d \geq 1.$$

This class consists of models nested within one another, where 'smaller' models can be obtained from the 'largest' model by setting specific parameters in the 'largest' model equal to specific values. For each $t \in [0, T]$, an observation Y is recorded and all observations are assumed to be independent normally distributed with the same variance, say $\sigma^2 > 0$, and the expectation of Y observed at t is given by

$$E[Y] = \eta(t, \theta)$$

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where $\eta(t, \theta)$ is one of the functions (2.3)-(2.6). In what is to follow, we suppress the p-dimensional parameter θ in $\eta(t, \theta)$ for simplicity when there is no confusion, and similarly for other notation such as $f(t, \theta)$ and $g(\theta)$ defined below.

In toxicological studies with a continuous outcome, the benchmark response is usually expressed in terms of a critical effect size (CES). This is the amount that we expect the percent change in the average level of the outcome compared with the background noise. In practice, CES is user-specified and traditionally set equal to 0.05 or 0.10. For a given mean response $\eta(t)$ and a user-selected CES, the critical effective dose CED is calculated from

$$\mathrm{CES} = -\frac{\eta(\mathrm{CED}) - \eta(0)}{\eta(0)}$$

if $\eta(t)$ is a decreasing function. All our functions $\eta(t)$ defined in (2.3)-(2.6) are decreasing.

The parameters in the above models may or may not all have meaningful interpretations, but frequently a re-parametrization of the mean function or working with a function of the model parameters has a practical meaning. By inverting the above functions, such as the mean function in (2.6), it is straightforward to show that

$$\mathrm{CED} = \mathrm{CED}(b, d, c) = \left(-\frac{\ln \frac{c-1+\mathrm{CES}}{c-1}}{b} \right)^{\frac{1}{d}}.$$

The corresponding expressions for CED for other models can be directly deduced by setting $c = 0$ for models (2.3) and (2.4) and by setting $d = 1$ for models (2.3) and (2.5). Thus the CED is the dose that results in a percent change in the mean response relative to the background noise and the magnitude of the anticipated change is specified by the critical effect size (CES).

To estimate the confidence interval for CED for a specific model using design ξ, one uses the delta method to obtain its asymptotic variance and then find a design to minimize it. Specifically, we have

$$\text{Var}(\widehat{\text{CED}}) \approx \frac{\sigma^2}{N} \Phi(\xi)$$

where

$$\Phi(\xi) = g^T(\theta) M^-(\xi, \theta) g(\theta), \quad \text{and} \quad g(\theta) = \frac{\partial \, \text{CED}}{\partial \theta}.$$

For the vector of the parameters $\theta = (a, b, d, c)^T$ **the local** CED**-optimal design** minimizes the function $\Phi(\xi)$ by choice of the design ξ. The matrix $M(\xi, \theta)$ in the above expressions is the information matrix for the specific model $\eta(t)$ using an arbitrary design ξ and $M^-(\xi, \theta)$ is a generalized inverse of $M(\xi, \theta)$. We call a design nonsingular if its information is nonsingular; otherwise it is a singular design. For a specific model η, let $f(t, \theta) = \frac{\partial \eta(t, \theta)}{\partial \theta}$ and recall that the information matrix is given by

$$M(\xi, \theta) = \int_0^T f(t, \theta) f^T(t, \theta) d\xi(t).$$

The corresponding regression vectors $f(t, \theta)$ for different models are

$$
\begin{aligned}
f^{(2.3)}(t, \theta) &= f(t, a, b) = (e^{-bx}, -ate^{-bt})^T, \\
f^{(2.4)}(t, \theta) &= f(t, a, b, d) = (e^{-bt^d}, -at^d e^{-bt^d}, -abt^d \ln(t) e^{-bt^d})^T, \\
f^{(2.5)}(t, \theta) &= f(t, a, b, c) = (c - (c-1)e^{-tb}, a(c-1)te^{-tb}, a(1 - e^{-tb}))^T, \\
f^{(2.6)}(t, \theta) &= (c - (c-1)e^{-bt^d}, a(c-1)t^d e^{-bt^d}, a(c-1)t^d \ln(t) be^{-bt^d}, a(1 - e^{-bt^d}))^T.
\end{aligned}
$$

and the corresponding vectors $g(\theta)$ are

$$g^{(2.3)}(\theta) = g(a,b) = \left(0, \frac{\ln(1-\text{CES})}{b^2}\right)^T$$

$$g^{(2.4)}(\theta) = \left(0, -\frac{1}{db}\left(-\frac{\ln(1-\text{CES})}{b}\right)^{\frac{1}{d}}, -\frac{1}{d^2}\left(-\frac{\ln(1-\text{CES})}{b}\right)^{\frac{1}{d}}\ln\left(-\frac{\ln(1-\text{CES})}{b}\right)\right)^T$$

$$g^{(2.5)}(\theta) = \left(0, \frac{\ln\frac{c-1+\text{CES}}{c-1}}{b^2}, \frac{1-\frac{c-1+\text{CES}}{c-1}}{b(c-1+\text{CES})}\right)^T,$$

$$g^{(2.6)}(\theta) = \left(0, -\frac{1}{db}\left(-\frac{\ln\frac{c-1+\text{CES}}{c-1}}{b}\right)^{\frac{1}{d}}, -\frac{1}{d^2}\left(-\frac{\ln\frac{c-1+\text{CES}}{c-1}}{b}\right)^{\frac{1}{d}}\ln\left(-\frac{\ln\frac{c-1+\text{CES}}{c-1}}{b}\right),\right.$$

$$\left.\left(-\frac{\ln\frac{c-1+\text{CES}}{c-1}}{b}\right)^{\frac{1}{d}}\frac{1-\frac{c-1+\text{CES}}{c-1}}{d(c-1+\text{CES})\ln\frac{c-1+\text{CES}}{c-1}}\right)^T.$$

The next five technical results provide analytical descriptions and properties of local CED-optimal design for each model. The first one shows that local CED-optimal design does not depend on the value of the parameter a, and the next four results describe the structure of the local CED-optimal designs for the four nonlinear models (2.3)-(2.6). We provide an illustrative proof of our results for model (2.3) only; the arguments for the other models are similar. For our class of models, the results also show how optimal design for a particular design setting can be deduced from another design setting by only considering values of b and T. Our technical justifications use a celebrated geometric result called Elfving's theorem which is widely discussed in design monographs, such as in Pázman (1986, p.71) or Pukelsheim (1993, p.50). We provide only proofs for Lemma 2.1 and 2.2; the rest are similar.

Lemma 2.1 *A local CED-optimal design does not depend on a.*

Proof. The statement follows from the fact that an optimality function have a form

$$\Phi(\xi, \theta) = a^{2-2p}\Phi(\xi, \tilde{\theta})$$

where $(a, \tilde{\theta}^T) = \theta^T \in \mathbb{R}^p$ and p is the number of parameters.

Lemma 2.2 *Let $u^* \approx 1.278$ be a unique solution of equation $e^{-u} = u - 1$. For model (2.3) a local CED-optimal design does not depend on a and CES and is given by*

$$\left\{ 0, u^*/b; \frac{e^{-u^*}}{1 + e^{-u^*}}, \frac{1}{1 + e^{-u^*}} \right\}$$

if $T > u^*/b$; otherwise it is given by

$$\left\{ 0, T; \frac{e^{-bT}}{1 + e^{-bT}}, \frac{1}{1 + e^{-bT}} \right\}.$$

Proof. By Elfving's theorem, there exists a representation

(2.7)
$$vg = w_1^* f(t_1^*) - w_2^* f(t_2^*)$$

for some $v \in \mathbb{R}$ and $g^T M^-(\xi^*) g = 1/v^2$. Moreover, points of optimal design lie on the boundary of Elfving set. Thus, $t_1^* = 0$. For large enough values of T, the point $f(t_2)$ belongs to the boundary if t_2 is small; otherwise it does not. The crucial value of t_2 is a solution of the equation

$$\frac{f_2'(t_2)}{f_1'(t_2)} = \frac{f_2(t_2) + f_2(0)}{f_1(t_2) + f_1(0)}.$$

Straightforward calculation shows that $t_2^* = u^*/b$. From the equation for the first coordinate of (2.7), we determine directly the weights w_1^* and w_2^* for the optimal design.

Lemma 2.3 *For model (2.4), the local CED-optimal design has one of three possible forms. It is either given by 2-point singular design*

$$\left\{ 0, CED; \frac{1 - CES}{2 - CES}, \frac{1}{2 - CES} \right\}$$

if the parameter b is small enough, or it has the form

$$\{0, t_2^*, t_3^*; w_1^*, w_2^*, w_3^*\}$$

if $t_3^* < T$; otherwise it takes the form

$$\{0, t_2^*, T; w_1^*, w_2^*, w_3^*\}.$$

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Moreover, for 3-point optimal designs, we have

$$t_i^*(b, d, T) = Tt_i^*(bT^d, d, 1), \quad w_i^*(b, d, T) = w_i^*(bT^d, d, 1).$$

$$t_i^*(b, 1, 1) = (t_i^*(b, d, 1))^d, \quad w_i^*(b, 1, 1) = w_i^*(b, d, 1).$$

For a 3-point optimal design, Elfving theorem implies that the weights of the optimal design are solutions of the equation

$$(g \vdots f(t_3^*) - f(0) \vdots f(t_3^*) + f(t_2))(v, w_1, w_2)^T = f(t_3^*)$$

with $t_2 = t_2^*$ and 2nd point t_2^* is a solution of $\partial v / \partial t_2 = 0$ where

$$v = v(t_2) = \frac{\det(f(t_3^*) \vdots f(t_3^*) - f(0) \vdots f(t_3^*) + f(t_2))}{\det(g \vdots f(t_3^*) - f(0) \vdots f(t_3^*) + f(t_2))}.$$

There is no explicit solution for model (2.4) but is an explicit solution for model (2.5).

Lemma 2.4 *For model* (2.5)*, the local CED-optimal design has one of two forms. It is either given by 2-point singular design*

$$\left\{ 0, CED; \frac{1 - CES}{2 - CES}, \frac{1}{2 - CES} \right\}$$

if the parameter b is small enough or c is large enough; otherwise, it has the following form

$$\{0, t_2^*, T; w_1^*, w_2^*, w_3^*\},$$

where

$$t_2^* = t_2^*(b, c, T) = \frac{1 - (1 + bT)e^{-bT}}{b(1 - e^{-bT})}.$$

Moreover, for 3-point optimal designs, we have

$$t_i^*(b, c, T) = Tt_i^*(bT, c, 1), \quad w_i^*(b, c, T) = w_i^*(bT, c, 1),$$

points $t_i^*(b, c, T)$ do not depend on c.

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Lemma 2.5 *For model* (2.6), *the local CED-optimal design has one of two form. It is either a singular 2-point design given by*

$$\left\{0, CED; \frac{1 - CES}{2 - CES}, \frac{1}{2 - CES}\right\}$$

if the parameter b is small enough or c is large enough; otherwise it has the form

$$\{0, t_2^*, t_3^*, T; w_1^*, w_2^*, w_3^*, w_4^*\}.$$

Moreover, for 4-point optimal designs, we have

$$t_i^*(b, d, c, T) = T t_i^*(bT^d, d, c, 1), \quad w_i^*(b, d, c, T) = w_i^*(bT^d, d, c, 1),$$

$$t_i^*(b, 1, c, 1) = (t_i^*(b, d, c, 1))^d, \quad w_i^*(b, 1, c, 1) = w_i^*(b, d, c, 1)$$

and points $t_i^*(b, d, c, T)$ do not depend on c.

Tables 1 and 2 show local CED-optimal designs for each of the four models when $CES = 0.05$ and selected values for b and T. As is described in the above results, the local CED-optimal design may be a singular 2-point design for models (2.4)-(2.6) or a saturated design where the number of points equal to the number of model parameters.

3   Maximin CED-optimal Design and Compound Optimal Design

The local optimal design for estimating CED depends on the assumed model and the nominal values of the model parameters. When the nominal values are mis-specified, local optimal designs can lose substantial efficiency. This problem is further compounded when there is model uncertainty. This implies that local optimal designs while potentially useful as a starting point, are unlikely to be adequate for practical

Table 1: *Local CED-optimal designs for model* (2.3) *and for model* (2.6) *with* $d = 1$, $c = 0$ *on the design space* $[0, T]$ *for various values of the parameter* b *and* CES $= 0.05$.

		model (2.3)				model (2.6)							
T	b	t_1	t_2	w_1	w_2	t_1	t_2	t_3	t_4	w_1	w_2	w_3	w_4
1	0.1	0	1	0.475	0.525	0	0.513			0.487	0.513		
1	0.5	0	1	0.377	0.623	0	0.103			0.487	0.513		
1	1.0	0	1	0.269	0.731	0	0.113	0.596	1	0.388	0.479	0.097	0.036
5	0.1	0	5	0.377	0.623	0	0.513			0.487	0.513		
5	0.5	0	2.557	0.218	0.782	0	0.430	2.482	5	0.337	0.463	0.141	0.058
5	1.0	0	1.278	0.218	0.782	0	0.277	1.718	5	0.316	0.454	0.159	0.071

Table 2: *Local CED-optimal designs for model* (2.4) *with* $d = 1$ *and for model* (2.5) *with* $c = 0$ *on the design space* $[0, T]$ *for various values of the parameter* b *and* CES $= 0.05$.

		model (2.4)						model (2.5)					
T	b	t_1	t_2	t_3	w_1	w_2	w_3	t_1	t_2	t_3	w_1	w_2	w_3
1	0.1	0	0.513		0.487	0.513		0	0.513		0.487	0.513	
1	0.5	0	0.305	1	0.378	0.515	0.106	0	0.459	1	0.353	0.532	0.115
1	1.0	0	0.251	1	0.344	0.511	0.145	0	0.418	1	0.317	0.554	0.129
5	0.1	0	1.523	5	0.378	0.515	0.106	0	2.293	5	0.353	0.532	0.115
5	0.5	0	0.672	4.507	0.305	0.462	0.233	0	1.553	5	0.261	0.593	0.146
5	1.0	0	0.336	2.253	0.305	0.462	0.233	0	0.966	5	0.232	0.613	0.155

implementation. However, local optimal designs are useful as a first step to constructing more versatile and robust designs to model assumptions. We now discuss two

design strategies that utilize local optimal designs.

The maximin approach of designing a study provides an alternative that can be appealing [see Dette (1995), Müller (1995) and Müller and Pázman (1998) among others]. Procedurally, one first considers the efficiency of a design for estimating the CED relative to each of the models; among these relative efficiencies, the maximin CED-optimal design is the one that maximizes the minimum of these relative efficiencies. Technically, for a fixed θ we call the design that maximizes

$$\min\{\text{eff}_{\text{CED}}^{(2.3)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.4)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.5)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.6)}(\xi,\theta)\}$$

over all designs on the design space a **maximin CED-optimal design**, where CED-efficiency (for each model) is given by

$$\text{eff}_{\text{CED}}(\xi) = \frac{\min_{\tilde{\xi}} \Phi(\tilde{\xi})}{\Phi(\xi)}.$$

The last ratio is obviously a number between 0 and 1 and represents the reduction in sample size from use of the optimal design compared with using the design ξ for the same level of precision for estimating CED. For example if $\text{eff}_{\text{CED}}(\xi) = 0.5$, the design ξ needs to be replicated twice to obtain an CED estimate as accurate as the estimate from the local optimal design for estimating CED. As expected, maximin optimal designs are difficult to find and defy analytical description. In particular, no closed form formulae are available. They have to be found numerically and several maximin optimal designs are shown in Table 3 for selected values of b and T with their efficiencies relative to the local CED-optimal designs for models (2.3)-(2.6).

In practice, maximin CED-optimal designs are found by first maximizing the optimality criterion within the class of all 4-point designs on the given design space. This is because 4 points are required for CED estimation in all models (2.3)-(2.6). The optimization is performed with the NelderMead algorithm in the MATLAB package. After the optimal

4-point design is found, we next search for the optimal design within the class of all 5 points designs, and repeat the procedure, each time increase the number of points by unity, until no reduction in the criterion value is observed.

Table 3: *Maximin CED-optimal designs for models* (2.3)-(2.6) *on the design space* $[0, T]$ *for various values of the parameter* b *with* $d = 1$, $c = 0$ *and* CES = 0.05 *and their efficiencies.*

T	b	t_1	t_2	t_3	t_4	w_1	w_2	w_3	w_4	$\text{eff}_{\text{CED}}^{(2.3)}$	$\text{eff}_{\text{CED}}^{(2.4)}$	$\text{eff}_{\text{CED}}^{(2.5)}$	$\text{eff}_{\text{CED}}^{(2.6)}$
1	0.1	0		0.513	1	0.417		0.296	0.287	0.686	0.686	0.686	0.692
1	0.5	0	0.183	0.612	1	0.299	0.228	0.231	0.242	0.597	0.636	0.597	0.597
1	1.0	0	0.170	0.594	1	0.261	0.217	0.288	0.234	0.615	0.615	0.615	0.615
5	0.1	0	0.914	3.059	5	0.299	0.228	0.231	0.242	0.597	0.636	0.597	0.597
5	0.5	0	0.820	2.451	5	0.237	0.292	0.350	0.121	0.655	0.655	0.655	0.655
5	1.0	0	0.529	1.545	5	0.236	0.366	0.326	0.072	0.639	0.639	0.639	0.673

3.1   Multiple-objective Optimal Designs

In practice, designs have several objectives in mind. There are interests in estimating various parameters in the model and frequently not all are of equal interest. For instance in the Michaelis-Menten model, the Michaelis-Menten constant b is clearly of much greater interest than the constant a. This means that the design should provide much more accurate estimate for the parameter b (the primary objective) than the parameter a (secondary objective). In this case, one may require that the design deliver at least 90% efficiency for estimating b and subject to this constraint devote the rest of the resources to estimating a. This is an example of a constrained optimal design discussed seminally in Stigler (1971), Studden (1982) and Lee (1988), where

they considered homoscedastic polynomial models. Such optimal designs are easy to motivate and interpret but usually they are difficult to find. When the objectives can be expressed as convex functionals of the information matrix, Cook and Wong (1994) proposed finding dual-objective optimal designs indirectly by first finding compound optimal designs.

Given two convex optimality criteria, it is tempting to consider a convex combination of the two criteria, which is also convex. This implies that for each λ between 0 and 1, one can straightforwardly find the compound optimal design ξ_λ that minimizes the convex combination. We next construct the efficiency plot that graphs the two efficiencies of ξ_λ versus values of λ between 0 and 1. Cook and Wong (1994) showed that the efficiency of ξ_λ under the primary criterion is always monotonically increasing and the efficiency under the secondary criterion is always monotonically decreasing. This is not surprising because resources have to be compromised for attaining the dual objectives. The slopes in the plots show how competitive the two objectives are, with steep slopes representing that much of one type of efficiency has to be given up in exchange for attaining more of the other efficiency. To relate compound optimal designs to constrained optimal designs, one uses the efficiency plots to arrive at a meaningful choice of λ. Specifically, the desired constrained optimal design is found by first drawing a horizontal line across the plot at the desired efficiency level sought for under the primary criterion, say 90% as in the above illustration. We then note the value of λ, say λ^*, that corresponds to the point where this horizontal line meets the (increasing) efficiency plot for the primary criterion and conclude that ξ_{λ^*} is the sought constrained optimal design. Details and theoretical explanation, along with worked out examples and illustrative efficiency plots are given in Cook and Wong (1994), Zhu, Zeng and Wong (2000), and Zhu and Wong (2000). See also Imhof and Wong (2000) where efficiency plots were used to find maximin optimal design in nonlinear models.

The advantage of the setup just described is that the researcher is in full control; he or she decides which is the primary criterion and sets the efficiency level required under the first criterion. This represents the level he or she is willing to compromise on the two objectives. Clearly, this method also works when all criteria are concave instead of convex, in which case we seek to maximize rather than minimize each criterion. The procedure just outlined can also be extended to find multiple-objective optimal designs; in our problem at hand, we set design criteria for discriminating between each pair of models and for CED estimation. Specifically, for a fixed θ we call the design that maximizes

$$\min\{\text{eff}_{\text{CED}}^{(2.3)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.4)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.5)}(\xi,\theta), \text{eff}_{\text{CED}}^{(2.6)}(\xi,\theta),$$

$$\text{eff}^{(2.4)-(2.3)}(\xi,\theta), \text{eff}^{(2.5)-(2.3)}(\xi,\theta), \text{eff}^{(2.6)-(2.5)}(\xi,\theta), \text{eff}^{(2.6)-(2.4)}(\xi,\theta)\}$$

over all designs on the design space a **maximin compound design** where $\text{eff}^{(M_1)-(M_2)}(\xi,\theta)$ is an efficiency of design ξ for discrimination two models M_1 and M_2. Such designs are efficient for CED estimation and discrimination. Selected maximin optimal designs are shown in Table 4 and their efficiencies for estimating CED under each model and for discriminating between pairs of models are shown in Table 5.

Our proposed maximin compound designs still depend on nominal values of the parameters. One could incorporate this uncertainty of the nominal values in a compound optimality criterion. We refer the reader to Biedermann, Dette and Pepelyshev (2006), Dette et al. (2008) for details but note that our small scale investigation showed that maximin optimal designs are not sensitive to small changes in the nominal values of the parameters in the models considered here.

Table 4: *Maximin compound designs for models (2.3)-(2.6) on the design space $[0, T]$ for various values of the parameter b with $d = 1$, $c = 0$ and CES $= 0.05$.*

T	b	t_1	t_2	t_3	t_4	w_1	w_2	w_3	w_4
1	0.1	0	0.173	0.622	1	0.361	0.152	0.281	0.205
1	0.5	0	0.183	0.612	1	0.299	0.228	0.231	0.242
1	1.0	0	0.170	0.594	1	0.261	0.217	0.288	0.234
5	0.1	0	0.914	3.059	5	0.299	0.228	0.231	0.242
5	0.5	0	0.698	2.398	5	0.186	0.225	0.298	0.291
5	1.0	0	0.639	1.899	5	0.147	0.267	0.240	0.347

Table 5: *CED-Efficiencies and efficiencies of discrimination of maximin compound designs for models (2.3)-(2.6) on the design space $[0, T]$ for various values of the parameter b with $d = 1$, $c = 0$ and CES $= 0.05$.*

T	b	CED efficiency				efficiency of discrimination			
		(2.3)	(2.4)	(2.5)	(2.6)	(2.4)-(2.3)	(2.5)-(2.3)	(2.6)-(2.5)	(2.6)-(2.4)
		maximin compound design							
1	0.1	0.610	0.620	0.620	0.610	0.622	0.651	0.610	0.694
1	0.5	0.597	0.637	0.597	0.597	0.678	0.634	0.720	0.737
1	1.0	0.615	0.615	0.615	0.615	0.643	0.626	0.743	0.803
5	0.1	0.597	0.637	0.597	0.597	0.677	0.634	0.720	0.737
5	0.5	0.605	0.633	0.605	0.605	0.697	0.605	0.716	0.813
5	1.0	0.456	0.456	0.596	0.474	0.456	0.456	0.527	0.517

4 Discussion

This closing section has four purposes aimed at the practitioners. The first purpose is to address distributional assumption on the error terms; in particular we construct

optimal designs under the assumption of log-normality and compare results obtained under the normality assumption. Secondly, we discuss efficiencies of designs used by toxicologists relative to our proposed optimal designs. The third purpose is to evaluate how well our maximin optimal designs perform in practice using a small simulation study. The final purpose is to draw attention to our design web site where many types of optimal designs for several models can be generated under user-specified settings.

4.1   Distributional assumptions

Sometimes toxicologists prefer to assume the continuous outcomes are log-normally distributed, see for example, Slob (2002). The dose-response model is fitted on the log-scale, where both the model and the data are log-transformed. After fitting the model, the model and the data may be back-transformed to the original scale for purposes of plotting and interpretation. We now show how our method can be extended to accommodate the log-normality assumption to find optimal designs.

It suffices to note here that we now assume logarithm of different observations are independent with the same variance, say $\sigma^2 > 0$, and have expectation

$$E[\ln Y] = \ln \eta(t, \theta)$$

where $\eta(t, \theta)$ is one of the 4 functions listed at the beginning of Section 2. Proceeding as in Section 2, one obtains an expression for the CED and the asymptotic variance of the estimated CED. The key difference is that the information matrix is now given by

$$\tilde{M}(\xi, \theta) = \int_0^T \tilde{f}(t, \theta) \tilde{f}^T(t, \theta) d\xi(t)$$

where $\tilde{f}(t, \theta) = \frac{1}{\eta(t,\theta)} \frac{\partial \eta(t,\theta)}{\partial \theta}$.

It follows that the vector $\tilde{f}(t,\theta)$ for each model is now different from the one under normality assumption in Section 2. However, the vector $g(\theta)$ for each model remains the same whether we assume the errors are normally or log-normally distributed. The next few tables display selected maximin CED-optimal designs (Table 6), maximin compound optimal designs (Table 8) under log-normal assumption and their efficiencies. From Tables 3 and 6, we observe that the maximin optimal designs obtained under the normality and log-normality assumptions do not appear to be substantially different. The same is observed for maximin compound designs in Tables 4 and 7.

Table 6: *Maximin CED-optimal designs for models (2.3)-(2.6) with lognormality assumption on the design space $[0,T]$ for various values of the parameter b with $d=1$, $c=0$ and $\mathrm{CES}=0.05$ and their efficiencies.*

T	b	t_1	t_2	t_3	t_4	w_1	w_2	w_3	w_4	$\mathrm{eff}_{\mathrm{CED}}^{(2.3)}$	$\mathrm{eff}_{\mathrm{CED}}^{(2.4)}$	$\mathrm{eff}_{\mathrm{CED}}^{(2.5)}$	$\mathrm{eff}_{\mathrm{CED}}^{(2.6)}$
1	0.1	0		0.513	1	0.430		0.289	0.281	0.691	0.691	0.691	0.691
1	0.5	0	0.196	0.638	1	0.334	0.224	0.221	0.221	0.620	0.655	0.620	0.620
1	1.0	0	0.195	0.655	1	0.326	0.202	0.240	0.232	0.636	0.636	0.636	0.636
5	0.1	0	0.982	3.191	5	0.334	0.224	0.221	0.221	0.620	0.655	0.620	0.620
5	0.5	0	1.188	3.485	5	0.334	0.198	0.238	0.231	0.643	0.643	0.643	0.643
5	1.0	0	1.450	3.758	5	0.347	0.204	0.239	0.211	0.644	0.644	0.644	0.644

4.2 Efficiency of commonly used designs

Now we discuss efficiencies of designs described in Section 1 relative to our proposed optimal designs. Recall that these are some of the types of designs commonly used by toxicologists in practice. Specifically, designs and nominal values of parameters are

Table 7: *Maximin compound designs for models (2.3)-(2.6) with lognormality assumption on the design space $[0, T]$ for various values of the parameter b with $d = 1$, $c = 0$ and* CES $= 0.05$.

T	b	t_1	t_2	t_3	t_4	w_1	w_2	w_3	w_4
1	0.1	0	0.176	0.628	1	0.374	0.155	0.271	0.200
1	0.5	0	0.196	0.638	1	0.334	0.224	0.221	0.221
1	1.0	0	0.199	0.659	1	0.325	0.202	0.242	0.232
5	0.1	0	0.981	3.191	5	0.334	0.224	0.221	0.221
5	0.5	0	1.174	3.469	5	0.322	0.199	0.241	0.238
5	1.0	0	1.357	3.683	5	0.312	0.206	0.250	0.233

Table 8: *CED-Efficiencies and efficiencies of discrimination for compound optimal designs for models (2.3)-(2.6) on the design space $[0, T]$ with lognormality assumption and various values of the parameter b with $d = 1$, $c = 0$ and* CES $= 0.05$.

T	b	CED efficiency				efficiency of discrimination			
		(2.3)	(2.4)	(2.5)	(2.6)	(2.4)-(2.3)	(2.5)-(2.3)	(2.6)-(2.5)	(2.6)-(2.4)
		maximin compound design							
1	0.1	0.614	0.622	0.627	0.614	0.628	0.648	0.614	0.688
1	0.5	0.620	0.655	0.620	0.620	0.675	0.628	0.721	0.727
1	1.0	0.636	0.636	0.636	0.636	0.645	0.636	0.707	0.753
5	0.1	0.620	0.655	0.620	0.620	0.675	0.628	0.722	0.727
5	0.5	0.643	0.643	0.643	0.643	0.649	0.643	0.678	0.730
5	1.0	0.640	0.640	0.640	0.640	0.644	0.640	0.662	0.729

Table 9: *CED-Efficiencies and efficiencies of discrimination for designs used by toxicologists for models (2.3)-(2.6) with lognormality assumption on the design space $[0, T]$ for various values of the parameter b, d and c and* CES $= 0.05$ *where M^* is the minimal efficiency of maximin compound design.*

				CED efficiency				efficiency of discrimination			
b	c	d	M^*	(2.3)	(2.4)	(2.5)	(2.6)	(2.4)-(2.3)	(2.5)-(2.3)	(2.6)-(2.5)	(2.6)-(2.4)
				design 0, 0.5, 1.5, 3, 4.5, 6, 7.5 (in 100mg/kg scale), $T = 7.5$							
0.04	1	0	0.597	0.492	0.504	0.563	0.361	0.558	0.579	0.594	0.666
0.06	1	0	0.616	0.492	0.517	0.570	0.419	0.558	0.579	0.594	0.666
0.09	1	0	0.628	0.492	0.526	0.574	0.464	0.558	0.579	0.594	0.665
				design 0, 0.5, 1.5, 4.5 (in 100mg/kg scale), $T = 7.5$							
0.04	1	0	0.597	0.217	0.375	0.185	0.551	0.245	0.057	0.126	0.032
0.06	1	0	0.616	0.217	0.348	0.165	0.497	0.245	0.053	0.121	0.028
0.09	1	0	0.628	0.217	0.331	0.150	0.375	0.245	0.047	0.114	0.024
				design 0, 0.5, 1.5, 4.5 (in 100mg/kg scale), $T = 4.5$							
0.04	1	0	0.607	0.602	0.526	0.398	0.548	0.681	0.498	0.370	0.294
0.06	1	0	0.596	0.602	0.566	0.406	0.517	0.681	0.493	0.364	0.286
0.09	1	0	0.612	0.602	0.590	0.406	0.490	0.681	0.485	0.356	0.274
				design 0, 1.0, 1.7, 2.8, 4.7, 7.8, 13, 22, 36, 60 , $T = 60$							
0.10	1	0	0.638	0.380	0.491	0.352	0.360	0.493	0.328	0.346	0.257
0.10	.5	0	0.613	0.380	0.440	0.352	0.478	0.488	0.328	0.613	0.640
0.04	.5	0	0.609	0.380	0.411	0.449	0.360	0.488	0.417	0.610	0.641
0.10	1	.9	0.614	0.380	0.491	0.417	0.319	0.493	0.529	0.551	0.265

taken from Woutersen et al. (2001) and Piersma et al. (2002). We list their various efficiencies in Table 9 relative to our optimal designs constructed under the log-normality assumption. Generally, the efficiencies of these designs are low for estimating CED or for discriminating between competing models in the stipulated class. They range from as low as 2% to mostly below 50%, and in a couple of instances about 67% for model discrimination. We see that in nearly all cases the minimal efficiency of the maximin compound design is greater than the efficiencies of the designs used by toxicologists. Additional calculation not shown here also reveals that the efficiencies of maximin compound designs are higher than the corresponding efficiencies of the used designs by at least on 10%. This means that any confidence interval for CED constructed from the maximin optimal design is at least on 10% shorter than those from the used designs. In many cases the improvement is even more substantial. We also compare performance of these used designs with optimal designs constructed under the normality assumption and the over trend is quite similar.

4.3   Performance of maximin optimal design in practice

All our optimal designs were found under a large sample assumption. These optimal designs minimize the asymptotic variance of the estimated CED obtained via the delta method. For this reason it is important to investigate the superiority of the optimal designs for sample size observed in practice. The purpose here is to briefly compare variances of the estimated CED from the maximin CED-optimal design and a design used by toxicologists in a real example with a relatively small sample size.

In Piersma et al. (2002), rats were prenatally exposed to diethylstilbestrol and the implemented design ξ_u had 6 animals in each of the 10 dose groups at 0, 1.0, 1.7, 2.8, 4.7, 7.8, 13, 22, 36 and 60 mg/kg body weight per day. This means that we have 60 observa-

tions on the design space $[0, 60]$. The maximin CED-optimal design ξ_{mm} for $b = 0.1$, $d = 1$, $c = 0$ has four dose levels and is given by $\{0, 5.2, 15.4, 60; 14/60, 21/60, 19/60, 6/60\}$.

We simulate data with $a = 1$ and $\sigma = 0.05$ and several values of parameters b, d and c with 1000 replicates in each simulation. In Table 10 we report simulated normalized variances of the estimated CED from the two designs under normality assumption. It is reassuring that we observe that in all the cases considered here, the variance obtained from the maximin optimal design ξ_{mm} is consistently smaller than the variance obtained from the design ξ_u used in practice. This implies that use of our proposed designs can save experimental cost for toxicologists and more importantly, in reducing the number of animals required in the study. In addition, unlike designs used by toxicologists, our designs are based on firm statistical considerations.

Table 10: *Simulated normalized variance of CED for several true values of parameters.*

b	d	c	$\mathrm{Var}(\widehat{CED})$ with ξ_{mm}	$\mathrm{Var}(\widehat{CED})$ with ξ_u
0.10	1.0	0.0	3.77	5.56
0.10	0.5	0.0	172.1	250.3
0.06	0.5	0.0	1821.0	3011.2
0.10	1.0	0.2	13.84	20.81
0.10	1.0	0.9	1850.4	3106.8

4.4 A design web site for practitioners

We conclude this paper with a reference to our web site where algorithms for generating optimal designs in this paper will be implemented very shortly. We believe that to facilitate use of optimal design ideas in practice, design tools should be readily available to practitioners. We thank the National Institute of General Medical Sci-

ences for funding the construction and maintenance of the site. This site is housed at http://www.optimal-design.org/ and visitors can generate a variety of optimal designs. We expect that algorithms used to generate the optimal designs in this paper will be available on this site shortly.

Presently, the site contains a list of models commonly used in the biological sciences, along with information and references on optimal design issues. The visitor selects an appropriate model, an optimality criterion and inputs parameters for the design problem. The generated design is displayed, and when appropriate, is also accompanied by a plot of the directional derivative of the optimality criterion. Depending on the features exhibited in this plot, we may or may not confirm the optimality of the generated design over all designs on the design space. This site also calculates efficiencies of user-supplied designs so that practitioners can easily compare their designs with the optimum and make an informed decision whether to stray away from the optimum and if so by how much. For space considerations, we omit further discussion and refer the reader to our web site. We hope that the site will promote use of optimal design ideas in practice.

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