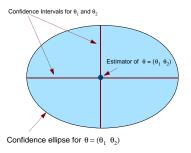
The criterion, introduced by Wald (1943), is

$$\Phi_D = \det(M^{-1}).$$

#### Properties:

- it minimises the generalised variance of the parameter estimator,
- it minimises the volume of the parameter confidence ellipsoid,
- ▶ it is invariant under linear transformations of the parameters,
- it is equivalent to G-optimality, which is given in the so-called Equivalence Theorem,
- it has at most p(p+1)/2 + 1 points of support (Carathéodory's Theorem).

Geometrical Interpretation - volume of confidence ellipsoid



A  $100(1-\alpha)\%$  confidence region for the parameters is

$$(\theta - \widehat{\theta})^{\mathrm{T}} M(\theta - \widehat{\theta}) \leq p s^2 F_{p,\nu,\alpha},$$

where  $s^2$  is an estimate of  $\sigma^2$ , and  $F_{p,\nu,\alpha}$  is the upper  $100\alpha\%$  point of the F distribution on p and  $\nu$  degrees of freedom.

The volume of a p-dimensional ellipsoid is proportional to  $\{\det(M^{-1})\}^{1/2}$ .

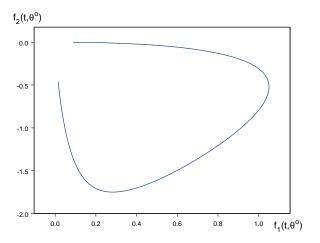
Geometrical Interpretation - design locus

Locally optimum designs for non-linear models with p parameters usually have p support points. Then the weights are all equal to 1/p.

The **design locus** is derived on the basis that the volume of a simplex in  $\mathbb{R}^p$ , formed by p points  $x_i \in \mathbb{R}^p$  and the origin, is proportional to the determinant of the  $(p \times p)$ -dimensional matrix formed by these points.

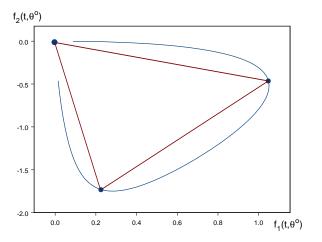
So, to maximise  $\det(M)$ , we find p points in the space of derivatives, which together with the origin, form a simplex of largest volume.

Geometrical Interpretation - design locus: pharmacokinetic model, p=2



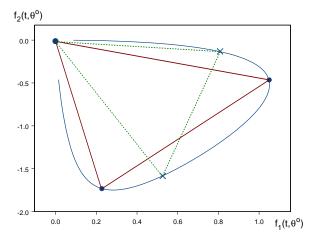
Design locus

Geometrical Interpretation - design locus: pharmacokinetic model, p=2



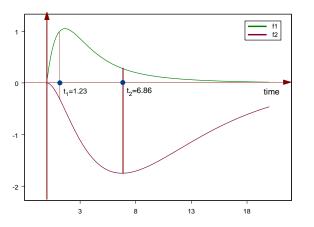
Design locus, optimum points and the simplex

Geometrical Interpretation - design locus: pharmacokinetic model, p=2



Design locus, optimum and non-optimum solutions

Geometrical Interpretation - parameter sensitivities



We find  $t_1$  and  $t_2$  such that  $det(X) = f_1(t_1)f_2(t_2) - f_2(t_1)f_1(t_2)$  is maximum.

The Equivalence Theorem

#### Kiefer and Wolfowitz (1960)

A design  $\xi^*$  is D-optimum if and only if it is G-optimum, that is, the following conditions are equivalent:

$$\det\{M^{-1}(\xi^*)\} = \min_{\xi} \det\{M^{-1}(\xi)\}$$

and

$$\max_{x} d(x, \xi^*) = \min_{\xi} \max_{x} d(x, \xi),$$

where  $d(x,\xi) = f(x)^{\mathrm{T}} M^{-1}(\xi) f(x)$  is the variance of prediction at a point x. The third equivalent condition says that

$$\max_{x} d(x, \xi^*) \le p,$$

where p is the number of parameters. Equality is achieved at the support points of  $\mathcal{E}^*$ .

The Equivalence Theorem, an Illustration

Let the model response be

$$\eta(x,\vartheta) = \vartheta_0 + \vartheta_1 x + \vartheta_2 x^2$$
 on  $[-1, 1]$ .

Then the D-optimum design is

$$\xi^* = \left\{ \begin{array}{rrr} -1 & 0 & 1\\ \frac{1}{3} & \frac{1}{3} & \frac{1}{3} \end{array} \right\}.$$

The design does not depend on n, but instead on the weights.

The information matrix can then be written as

$$M(\xi^{\star}, \vartheta^{o}) = X^{\mathrm{T}}WX = \begin{pmatrix} 1 & 1 & 1 \\ -1 & 0 & 1 \\ 1 & 0 & 1 \end{pmatrix} \times \begin{pmatrix} \frac{1}{3} & 0 & 0 \\ 0 & \frac{1}{3} & 0 \\ 0 & 0 & \frac{1}{3} \end{pmatrix} \times \begin{pmatrix} 1 & -1 & 1 \\ 1 & 0 & 0 \\ 1 & 1 & 1 \end{pmatrix}.$$

The Equivalence Theorem, an Illustration

Hence,

$$M = \frac{1}{3} \left( \begin{array}{ccc} 3 & 0 & 2 \\ 0 & 2 & 0 \\ 2 & 0 & 2 \end{array} \right)$$

and the variance function is

$$d(x,\xi^*) = f(x)^{\mathrm{T}} M^{-1} f(x)$$

$$= 3(1, x, x^2) \times \begin{pmatrix} 1 & 0 & -1 \\ 0 & 0.5 & 0 \\ -1 & 0 & 1.5 \end{pmatrix} \times \begin{pmatrix} 1 \\ x \\ x^2 \end{pmatrix}$$

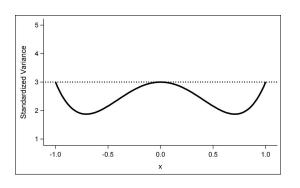
$$= 3 - 4.5x^2 + 4.5x^4.$$

Note that  $d(x, \xi^*) = 3$  at x = -1, 0, 1.

The Equivalence Theorem, an Illustration

#### Recall that

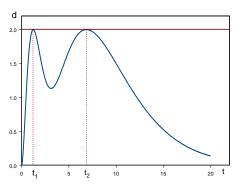
$$\xi^* = \left\{ \begin{array}{rrr} -1 & 0 & 1\\ \frac{1}{3} & \frac{1}{3} & \frac{1}{3} \end{array} \right\}.$$



The Equivalence Theorem - pharmacokinetic model

Here,

$$\xi^* = \left\{ \begin{array}{cc} 1.23 & 6.86 \\ \frac{1}{2} & \frac{1}{2} \end{array} \right\}.$$



## Example 4

Enzyme Kinetics Model, p=2

In a typical enzyme kinetics reaction, enzymes bind substrates and turn them into products. The binding step is reversible while the catalytic step is irreversible:

$$S + E \longleftrightarrow ES \to E + P$$
,

where S, E and P denote the substrate, enzyme and product, respectively.

#### Example 4

#### Enzyme Kinetics Model, p = 2

The reaction rate is represented by the Michaelis-Menten model

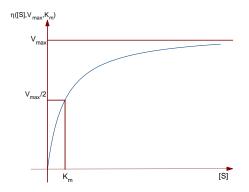
$$\eta([S]; V_{max}, K_m) = \frac{V_{max}[S]}{K_m + [S]},$$

where [S] is the concentration of the substrate, and  $V_{max}$  and  $K_m$  are the model parameters:

- $\triangleright$   $V_{max}$  denotes the maximum velocity of the reaction and
- $ightharpoonup K_m$  is the Michaelis-Menten constant, the value of [S] at which one half of the maximum velocity  $V_{max}$  is reached.

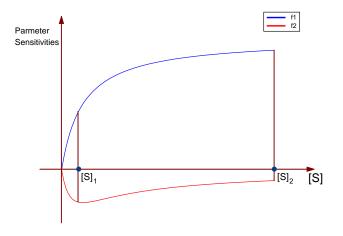
#### Example 4

Enzyme Kinetics Model, p = 2



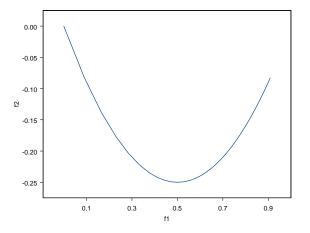
The Michaelis-Menten model response function  $\eta([S]; V_{max}, K_m)$  for the point priors  $V_{max}^o = 1$  and  $K_m^o = 1$ .

**Enzyme Kinetics Model**, p = 2, parameter sensitivities



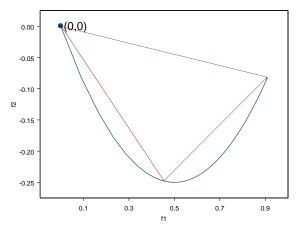
Note that  $f_1$  does not have a proper maximum; the largest value is at the boundary of the design region.

#### D optimality Enzyme Kinetics Model, p = 2, design locus



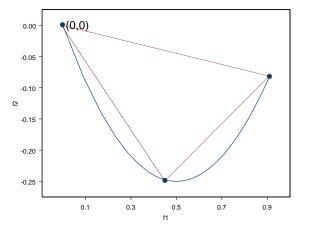
The design locus does not form a loop.

**Enzyme Kinetics Model**, p = 2, design locus



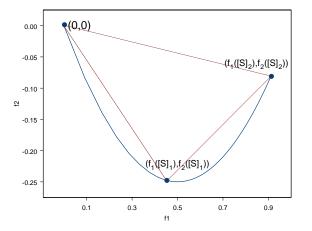
Design locus: one vertex must be at the end of the locus.

**Enzyme Kinetics Model**, p = 2, design locus



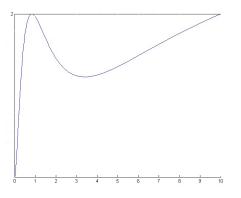
Design locus: the triangle of maximum area.

**Enzyme Kinetics Model**, p = 2, design locus



Design locus: optimum design points.

**Enzyme Kinetics Model**, p = 2, The Equivalence Theorem



The variance function has only one proper maximum; it also reaches p=2 at the boundary of the design region.

#### Atkinson and Bogacka (2002)

Suppose that

$$A \stackrel{k_1}{\rightarrow} B \stackrel{k_2}{\rightarrow} C.$$

Then the kinetic differential equations for [A], [B] and [C], the concentrations of the chemical compounds A, B and C as functions of time t, are

$$\frac{d[A]}{dt} = -k_1[A]^{\lambda_1},$$

$$\frac{d[B]}{dt} = k_1[A]^{\lambda_1} - k_2[B]^{\lambda_2},$$

$$\frac{d[C]}{dt} = k_2[B]^{\lambda_2}.$$

Interest is in estimation of the orders  $\lambda_1$  and  $\lambda_2$ , as well as of the rates  $k_1$  and  $k_2$ .

The first equation can be solved analytically to give the concentration of chemical A at time t as

$$[A] = \{1 - (1 - \lambda_1)k_1t\}^{1/(1-\lambda_1)}, \qquad \lambda_1, k_1, t \ge 0, \lambda_1 \ne 1,$$

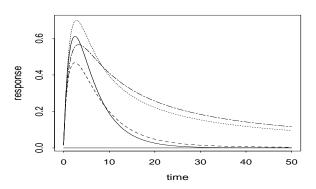
if it is assumed that the initial concentration of A is 1.

This gives the following differential equation for the concentration of the compound *B*:

$$\frac{d[B]}{dt} = k_1 \{1 - (1 - \lambda_1)k_1t\}^{\frac{\lambda_1}{1 - \lambda_1}} - k_2[B]^{\lambda_2},$$

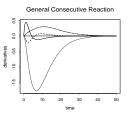
which has to be solved numerically.

#### General Consecutive Reaction



Concentration of *B*. Reading upwards at t = 20:  $(\lambda_1^o, \lambda_2^o) = (1, 1), (2, 1), (1, 2)$  and  $(2, 2), (k_1^o, k_2^o) = (0.7, 0.2)$ .

Parameter sensitivities



Parameter sensitivities as a function of time. Reading upwards at t=10:  $f_2, f_1, f_3$  and  $f_4$  for  $k_2, k_1, \lambda_1$  and  $\lambda_2$ , respectively. Here,  $(\lambda_1^o, \lambda_2^o) = (1, 1)$  and  $(k_1^o, k_2^o) = (0.7, 0.2)$ .

D-optimum designs

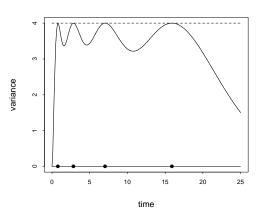
These designs were found by searching over the values of time, but with the weights held known at 0.25. The design region is  $\mathcal{T} = [0,50]$ .

Prior Rates and Orders	Time			
$(k_1^o,k_2^o,\lambda_1^o,\lambda_2^o)$	$t_1^*$	$t_2^*$	$t_3^*$	$t_4^*$
(0.7, 0.2, 1,1)	0.80	2.85	7.05	15.90
(0.7, 0.2, 2, 1)	0.51	2.36	7.30	18.26
(0.7, 0.2, 1, 2)	0.83	2.91	8.05	40.39
(0.7, 0.2, 2, 2)	0.57	2.65	9.68	50.00

Table 1. D-optimum designs for both rate and order. The weights are 0.25 at each design point.

D-optimum designs

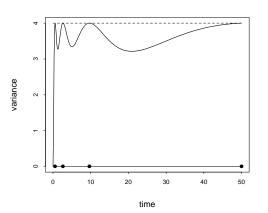
$$A -> B -> C$$
: lambda = (1,1)



The variance of prediction  $d(t, \xi^*, \vartheta)$  for prior  $(k_1^o, k_2^o, \lambda_1^o, \lambda_2^o) = (0.7, 0.2, 1, 1).$ 

D-optimum designs

$$A -> B -> C$$
: lambda = (2,2)



The variance of prediction  $d(t, \xi^*, \vartheta)$  for prior  $(k_1^o, k_2^o, \lambda_1^o, \lambda_2^o) = (0.7, 0.2, 2, 2).$