



UPPSALA
UNIVERSITET

Different approximations and methods for calculating the FIM and their consequences

Joakim Nyberg, Joakim Ringblom,
Mats O. Karlsson, Andrew C. Hooker

Division of Pharmacokinetics and Drug Therapy

Department of Pharmaceutical Biosciences

Uppsala University

Sweden

2008-06-23



Background

Several software are available in the Population Optimal Design area and they all have different options for how to calculate the FIM and also how to include residual variability in the model.

How can this effect the optimal design?



Introduction

- The Population Fisher Information Matrix (FIM) can be very time consuming to optimize over => The reduced FIM is quite often used, what consequences does that have on the optimization?
- The residual variance of a model is often proportional + additive (slope+const), how will different implementations of the variance model affect the optimal design?
- The FIM is often parameterized differently in terms of SD or variances, consequences?!



Full vs. Reduced FIM - Example

$$\begin{array}{c} \theta_{CL} \quad \theta_V \quad \omega_{CL}^2 \quad \omega_V^2 \\ \\ FIM_{Full}^{-1} = \begin{pmatrix} \theta_{CL} & \theta_{CL-V} & \theta_{CL-\omega_{CL}^2} & \theta_{CL-\omega_V^2} \\ \theta_{CL-V} & \theta_V & \theta_{V-\omega_{CL}^2} & \theta_{V-\omega_V^2} \\ \theta_{CL-\omega_{CL}^2} & \theta_{V-\omega_{CL}^2} & \omega_{CL}^2 & \omega_{CL-V}^2 \\ \theta_{CL-\omega_V^2} & \theta_{V-\omega_V^2} & \omega_{CL-V}^2 & \omega_V^2 \end{pmatrix} \\ \\ \theta_{CL} \quad \theta_V \quad \omega_{CL}^2 \quad \omega_V^2 \\ \\ FIM_{Reduced}^{-1} = \begin{pmatrix} \theta_{CL} & \theta_{CL-V} & & \\ \theta_{CL-V} & \theta_V & & \\ & & \omega_{CL}^2 & \omega_{CL-V}^2 \\ & & \omega_{CL-V}^2 & \omega_V^2 \end{pmatrix} \end{array}$$

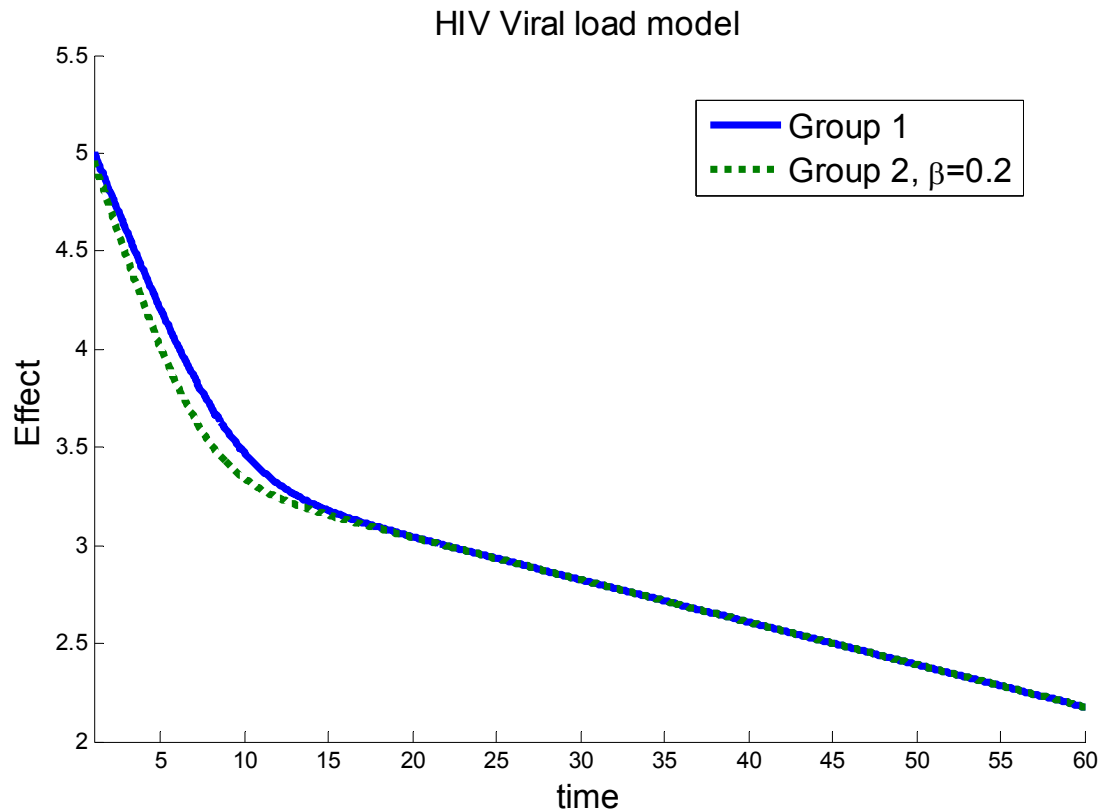


Different models investigated with Full vs. Reduced FIM

- First order absorption model
- Bateman function
- Biorhythm model
- Derendorf surge model
- Enterohepatic recirculation model
- IV PK/Emax PD model
- Lag-time model
- Michalis-Menten elimination model
- Pool-tolerance model
- Transit compartment absorption model
- Viral load model



HIV Viral load



$$E_{group1} = \log_{10}\left(e^{P_1} \cdot e^{-e^{d_1 \cdot t}} + e^{P_2} \cdot e^{-e^{d_2 \cdot t}}\right)$$

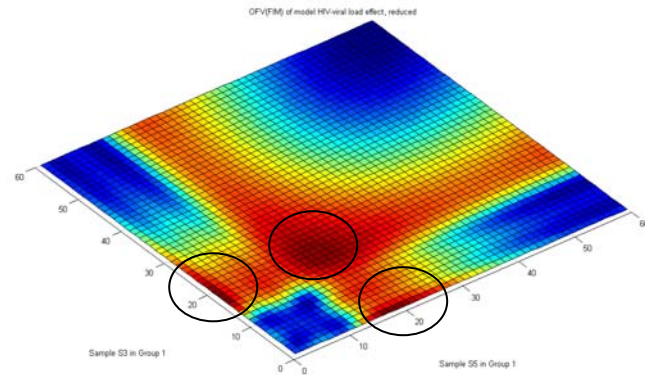
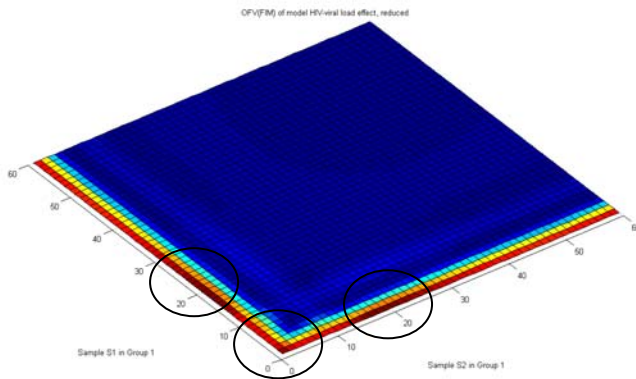
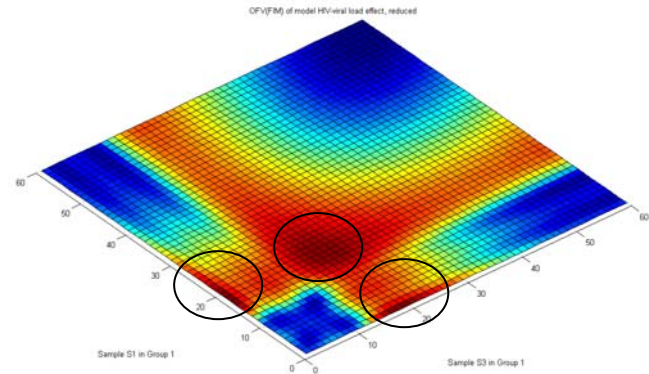
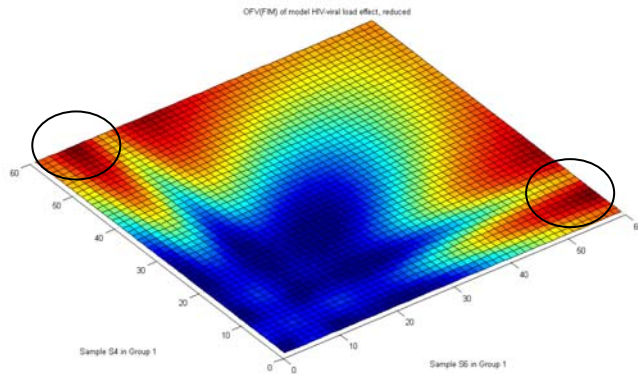
$$E_{group2} = \log_{10}\left(e^{P_1} \cdot e^{-e^{(d_1 + \beta) \cdot t}} + e^{P_2} \cdot e^{-e^{d_2 \cdot t}}\right)$$

Retout et al. *Statist. Med.* 2007; **26**:5162–5179. Design in nonlinear mixed effects models: Optimization using the Fedorov Wynn algorithm and power of the Wald test for binary covariates. 6



HIV Viral load model

Reduced FIM

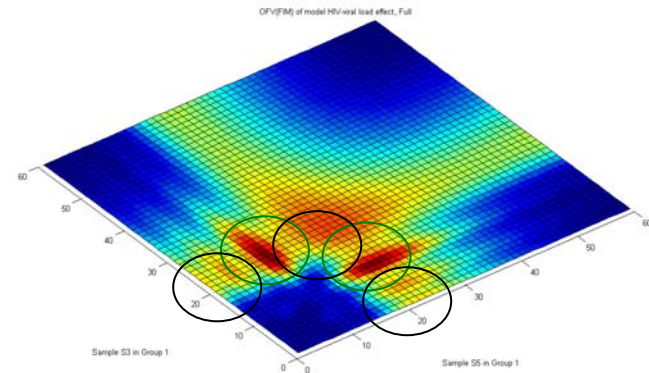
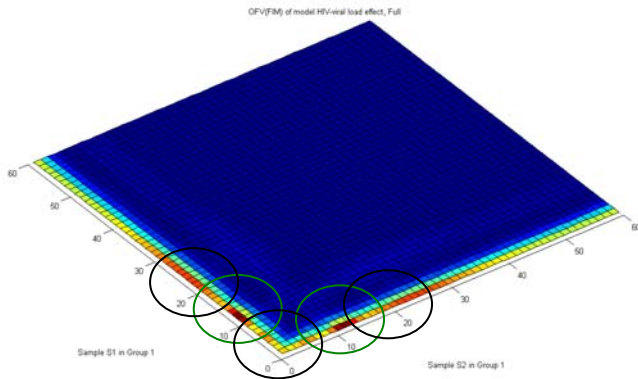
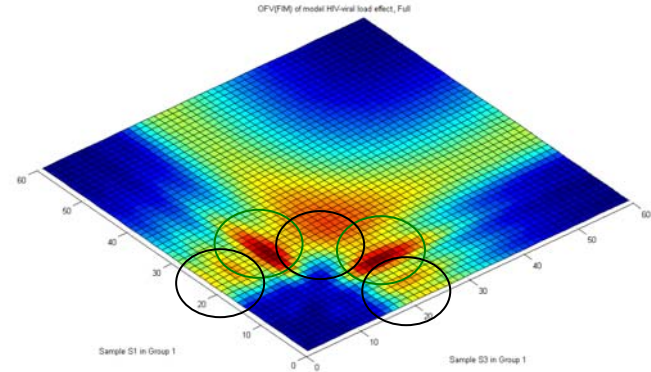
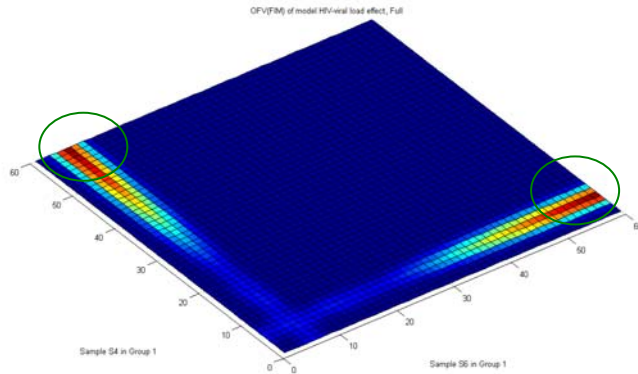


○ = optimal value



HIV Viral load model

Full FIM

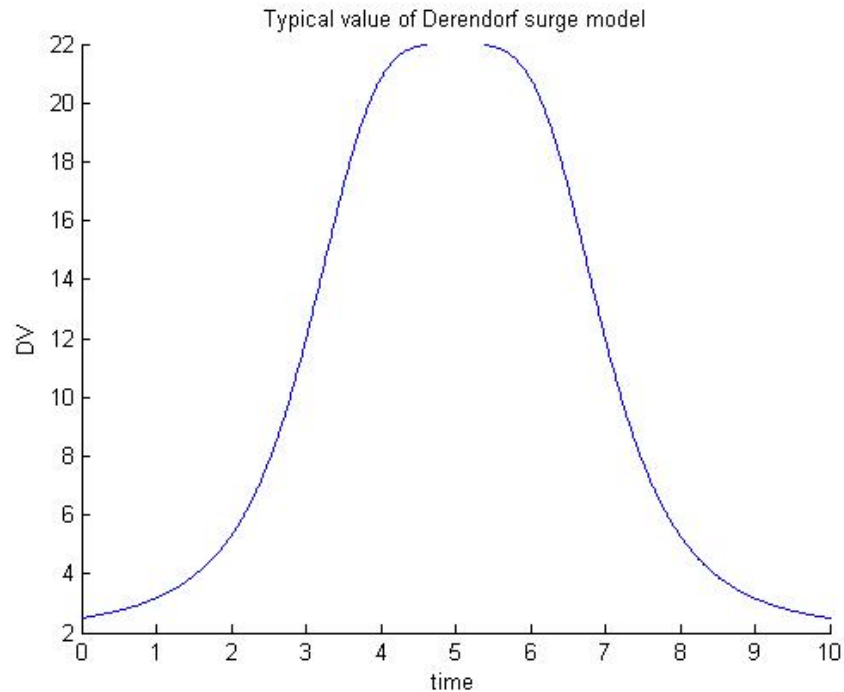


○ = optimal value



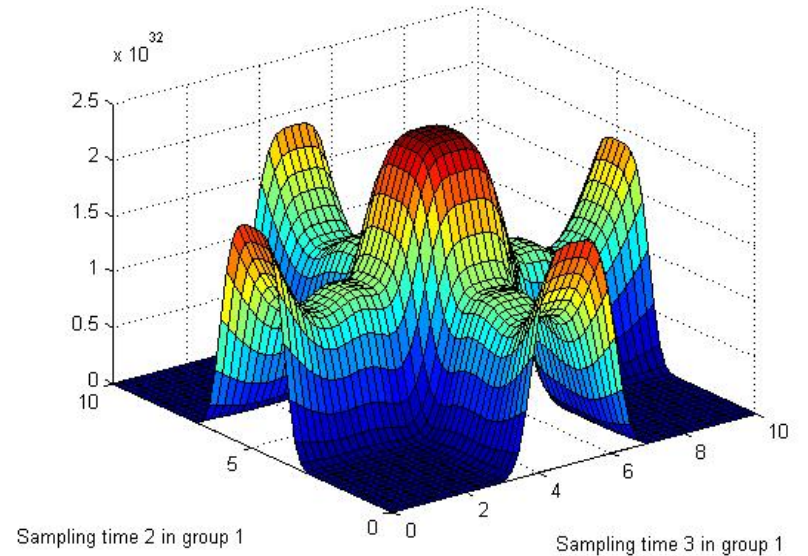
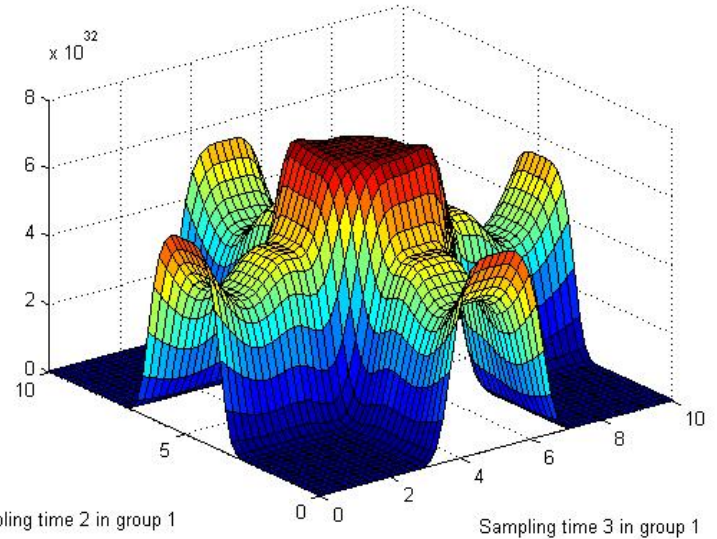
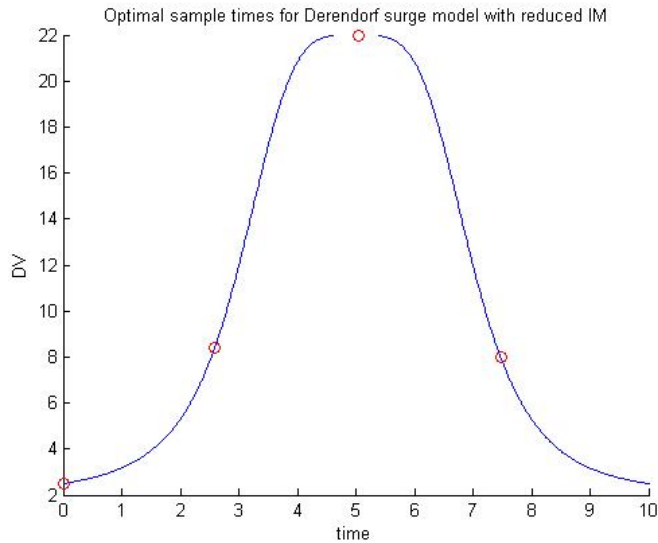
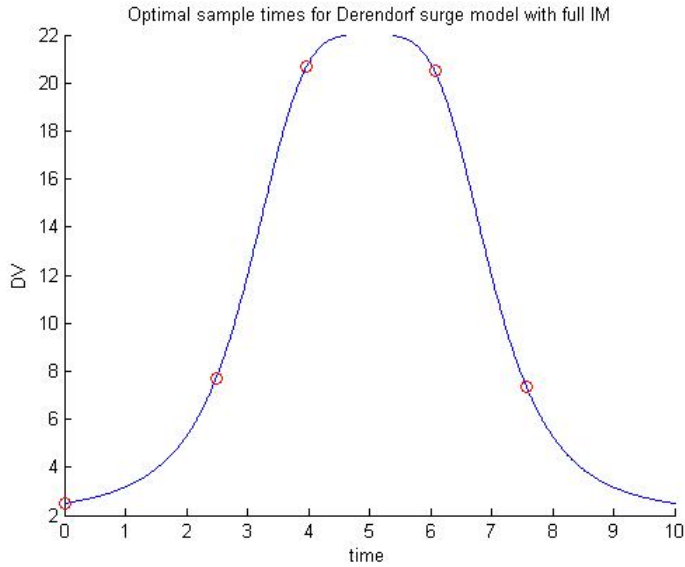
Derendorf surge model

$$y = R_0 \cdot \left(1 + \frac{SA}{\left(\frac{t - T_0}{SW} \right)^N + 1} \right)$$





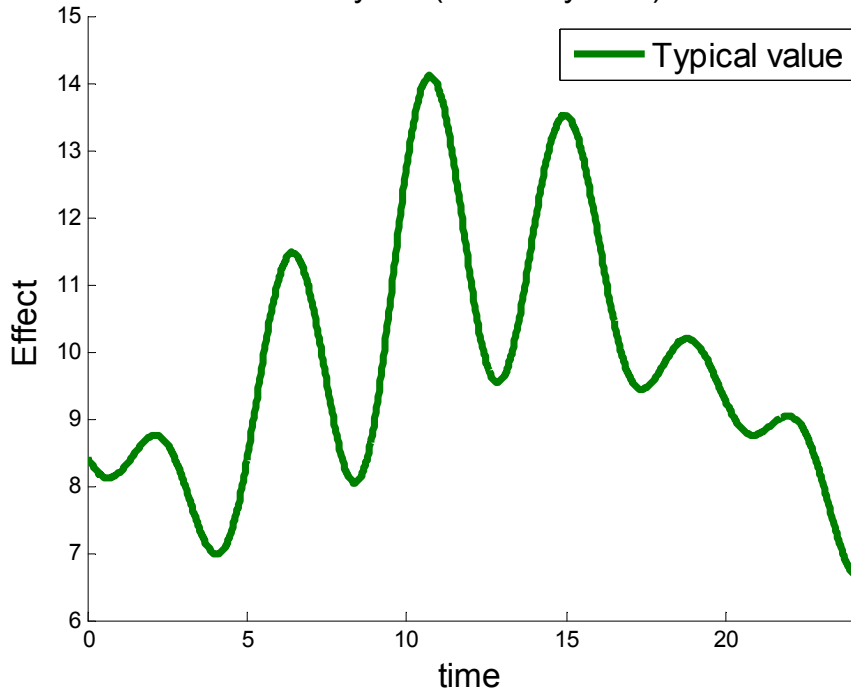
Derendorf surge model-OD



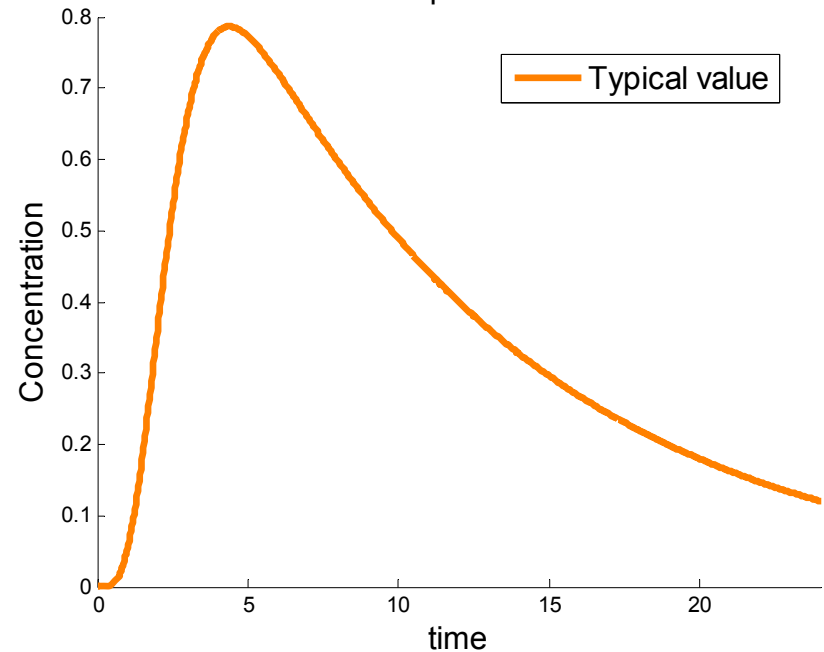


Biorhythm & Transit compartment

Biorhythm (3 cos rhythms)



Transit compartment model



$$A = \frac{k_{in}}{k_{out}} + \frac{amp \cdot k_{in}}{k_{out}^2 + \left(\frac{2\pi}{per}\right)^2} \cdot \left(\begin{array}{l} k_{out} \cdot \cos\left((t - peak) \cdot \frac{2\pi}{per}\right) + \\ \frac{2\pi}{per} \cdot \sin\left((t - peak) \cdot \frac{2\pi}{per}\right) \end{array} \right)$$

$$\frac{\partial A_{abs}}{\partial t} = e^{A_n} \cdot ((n+1) \cdot MTT) - k_a \cdot A_{abs}$$

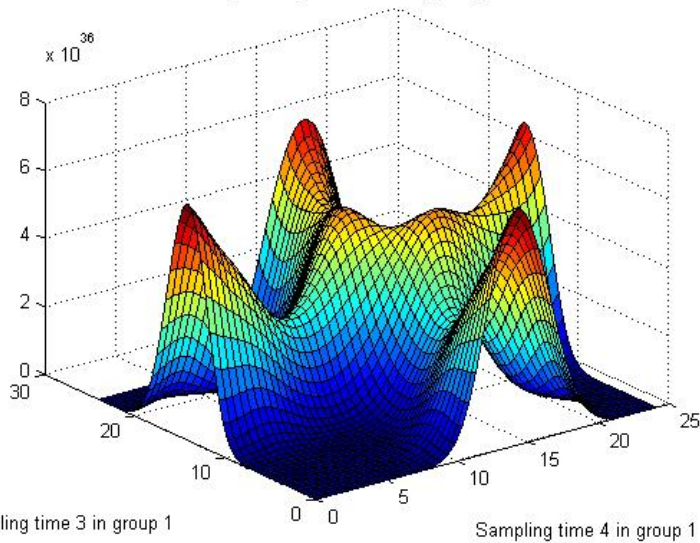
$$\frac{\partial A_{central}}{\partial t} = k_a \cdot A_{abs} - \frac{CL}{V} \cdot A_{central}$$



Biorhythm results

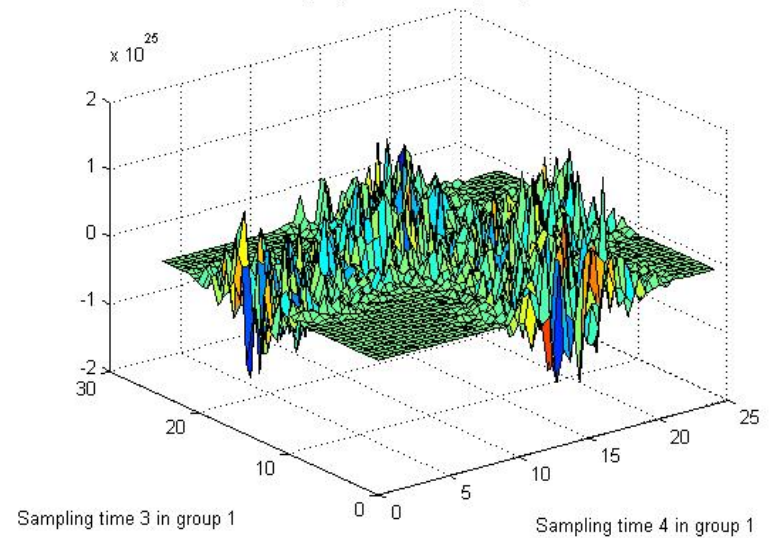
Full

OFV(full FIM) of model Biological rythm



Reduced

OFV(FIM) of model Biological rythm



- Singular FIM -> Hard/Impossible to optimize on!
- Similar for Transit compartment model



Why singularity for reduced?

Biorhythm:

Rows of $M_1 = \frac{\partial f(\bar{x}_i, \Theta)}{\partial \theta}$ are linearly dependent

Transit model:

Rows of $M_3 = \frac{\partial \text{vec}(\text{Var}(\bar{y}_i))}{\partial \omega^2}, \frac{\partial \text{vec}(\text{Var}(\bar{y}_i))}{\partial \sigma^2}$ are linearly dependent



UPPSALA
UNIVERSITET

Similar optimal designs with reduced FIM

- First order absorption model
- Bateman function model
- Enterohepatic recirculation model
- IV PK/Emax PD model
- Lag-time model
- Michaelis-Menten elimination model
- Pool-tolerance model



Residual Variability models

General model

$$\vec{y}_i = \mathbf{f}(\vec{\theta}_i, \vec{x}_i) + \mathbf{h}(\vec{\theta}_i, \vec{x}_i, \vec{\varepsilon}_i) \quad \vec{\varepsilon}_i \sim N(0, \Sigma) \quad \vec{\eta}_i \sim N(0, \Omega)$$

Additive + proportional residual variability model

$$\sigma_{add}^2 + \sigma_{prop}^2 \cdot f(\vec{\theta}, \vec{x})^2 \quad \vec{\varepsilon}_i \sim N(0, \Sigma)$$

Additive + proportional residual variability model

$$\left(\sigma_{add} + \sigma_{prop} \cdot f(\vec{\theta}, \vec{x})\right)^2 \quad \vec{\varepsilon}_i \sim N(0, \Sigma)$$



General Variance

General model

$$\vec{y}_i = f(\vec{\theta}_i, \vec{x}_i) + h(\vec{\theta}_i, \vec{x}_i, \vec{\varepsilon}_i)$$

$$\vec{\varepsilon}_i \sim N(0, \Sigma)$$

$$\vec{\eta}_i \sim N(0, \Omega)$$

General variance

$$\text{var}(\vec{y}_i) = \mathbf{L} \cdot \Omega \cdot \mathbf{L}^T + \text{diag}(\mathbf{H} \cdot \Sigma \cdot \mathbf{H}^T)$$

$$\mathbf{L}_i(\vec{x}_i, \vec{\theta}_i) \equiv \left. \frac{\partial f}{\partial \vec{\eta}_i}(\vec{x}_i, \vec{\theta}_i) \right|_{\vec{\eta}_i=0}$$

$$\mathbf{H}_i(\vec{x}_i, \vec{\theta}_i) \equiv \left. \frac{\partial h}{\partial \vec{\varepsilon}_i}(\vec{x}_i, \vec{\theta}_i, \vec{\varepsilon}_i) \right|_{\vec{\eta}_i=0, \vec{\varepsilon}_i=0}$$



General Variance applied to different residual models

$$\sigma_{add}^2 + \sigma_{prop}^2 \cdot f(\vec{\theta}, \vec{x})^2$$

$$\text{var}(\vec{y}_i) = \text{diag}(H \cdot \Sigma \cdot H^T) = \text{diag} \left(\begin{pmatrix} 1 & f(t_1, \vec{\theta}) \\ 1 & f(t_2, \vec{\theta}) \end{pmatrix} \cdot \begin{pmatrix} \sigma_{add}^2 & 0 \\ 0 & \sigma_{prop}^2 \end{pmatrix} \cdot \begin{pmatrix} 1 & f(t_1, \vec{\theta}) \\ 1 & f(t_2, \vec{\theta}) \end{pmatrix}^T \right) =$$
$$\begin{pmatrix} \sigma_{add}^2 + \sigma_{prop}^2 \cdot f(t_1)^2 & 0 \\ 0 & \sigma_{add}^2 + \sigma_{prop}^2 \cdot f(t_2)^2 \end{pmatrix}$$

$$\left(\sigma_{add} + \sigma_{prop} \cdot f(\vec{\theta}, \vec{x}) \right)^2$$

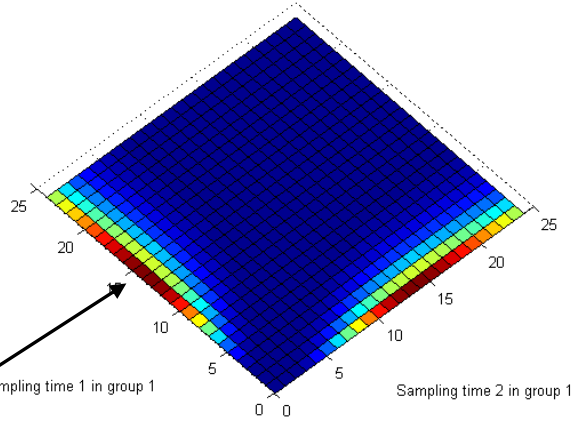
$$\text{var}(\vec{y}_i) = \text{diag}(H \cdot \Sigma \cdot H^T) = \text{diag} \left(\begin{pmatrix} 1 & f(t_1, \vec{\theta}) \\ 1 & f(t_2, \vec{\theta}) \end{pmatrix} \cdot \begin{pmatrix} \sigma_{add} & \sigma_{add} \cdot \sigma_{prop} \\ \sigma_{add} \cdot \sigma_{prop} & \sigma_{prop}^2 \end{pmatrix} \cdot \begin{pmatrix} 1 & f(t_1, \vec{\theta}) \\ 1 & f(t_2, \vec{\theta}) \end{pmatrix}^T \right) =$$
$$\begin{pmatrix} \left(\sigma_{add} + \sigma_{prop} \cdot f(t_1) \right)^2 & 0 \\ 0 & \left(\sigma_{add} + \sigma_{prop} \cdot f(t_2) \right)^2 \end{pmatrix}$$



Implications of residual models

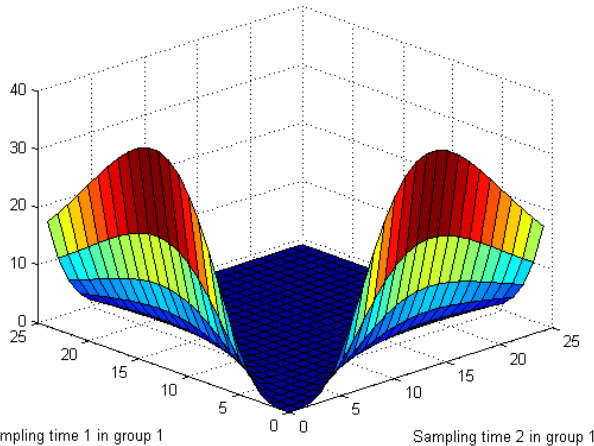
$$\sigma_{\text{add}}^2 + f^2 \cdot \sigma_{\text{prop}}^2$$

OFV(FIM) with $(a^2 + b^2 \cdot f^2)$ parameterization of residual error



Optimal

$$\text{OFV(FIM) with } (a^2 + b^2 \cdot f^2) \text{ parameterization of residual error}$$

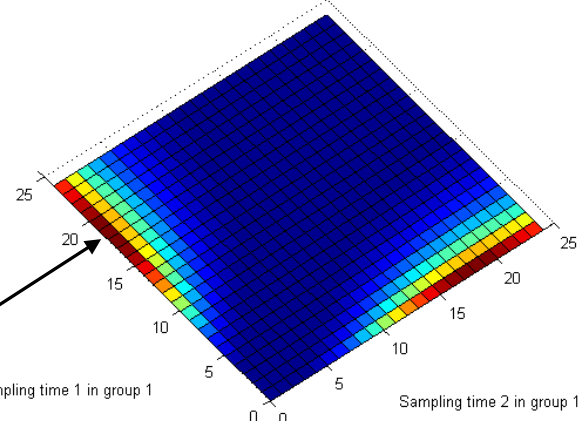


Sampling time 1 in group 1 Sampling time 2 in group 1

CV: $\theta_1 = 0.9$ $\sigma_{\text{add}}^2 = 1.6$
 $\theta_2 = 2.2$ $\sigma_{\text{prop}}^2 = 5.2$

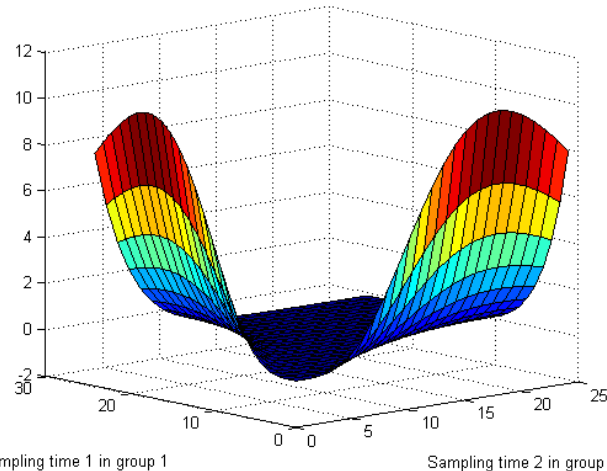
$$(\sigma_{\text{add}} + f \cdot \sigma_{\text{prop}})^2$$

OFV(FIM) with $(a + b \cdot f)^2$ parameterization of residual error



Optimal

$$\text{OFV(FIM) with } (a + b \cdot f)^2 \text{ parameterization of residual error}$$



Sampling time 1 in group 1 Sampling time 2 in group 1

CV: $\theta_1 = 1.3$ $\sigma_{\text{add}}^2 = 2.5$
 $\theta_2 = 2.8$ $\sigma_{\text{prop}}^2 = 5.9$



Different derivations of FIM, w.r.t. variance vs. stdev

Variance derivation

$$\frac{\partial \text{var}(\vec{y})}{\partial \sigma^2} = H \cdot H^T \quad \Longrightarrow \quad SE_{\sigma^2} = 2 \cdot SE_{\sigma}$$

Stdev derivation

$$\frac{\partial \text{var}(\vec{y})}{\partial \sigma} = 2 \cdot \sigma \cdot H \cdot H^T \quad \Longrightarrow \quad SE_{\sigma} = \frac{SE_{\sigma^2}}{2}$$

The difference in SE will reflect differences in |FIM| and vice versa



Conclusions

- It is important to increase the understanding of potential differences between software in OD.
- Different approximations and different residual models can clearly affect the optimal design and in some cases lead to different results.
- When comparing the expected uncertainty of an estimator, the residual variability model needs to be considered to get an accurate comparison.
- Allowing for a general error function and linearizing around it is the most flexible way to allow for all combinations of models and correlations.



Backup - Calculating FIM

$$\mathbf{FIM}(\Theta)_i = \begin{pmatrix} \mathbf{M}_{1i} & \mathbf{0} \\ \mathbf{M}_{2i} & \mathbf{M}_{3i} \end{pmatrix}^T \begin{pmatrix} \mathbf{Var}(\vec{y}_i)^{-1} & \mathbf{0} \\ \mathbf{0} & \mathbf{M}_{4i}^{-1} \end{pmatrix} \begin{pmatrix} \mathbf{M}_{1i} & \mathbf{0} \\ \mathbf{M}_{2i} & \mathbf{M}_{3i} \end{pmatrix}$$

$$\mathbf{M}_{1i} = \frac{\partial f(\vec{x}_i, \vec{\theta})}{\partial \vec{\theta}}$$

$$\mathbf{M}_{2i} = \frac{\partial \text{vec}(\mathbf{Var}(\vec{y}_i))}{\partial \vec{\theta}}$$

$$\mathbf{M}_{3i} = \left[\frac{\partial \text{vec}(\mathbf{Var}(\vec{y}_i))}{\partial \vec{\omega}^2}, \frac{\partial \text{vec}(\mathbf{Var}(\vec{y}_i))}{\partial \vec{\sigma}^2} \right]$$

$$\mathbf{M}_{4i} = 2 \cdot \mathbf{Var}(\vec{y}_i) \otimes \mathbf{Var}(\vec{y}_i)$$

$$\mathbf{Var}(\vec{y}_i) \approx \mathbf{L} \cdot \mathbf{\Omega} \cdot \mathbf{L}^T + \text{diag}(\mathbf{H} \cdot \mathbf{\Sigma} \cdot \mathbf{H}^T)$$