

### **Modelling transfer of dioxins from feed to eggs**

experimental data model building identification analysis

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### Modelling in three stages:

#### •Experimental data:

no data obtained, no knowledge gained <u>data obtained, knowledge gained</u>

#### •Model building:

PBPK, classical compartment, classical kinetic do not stick to paradigms

#### Identification analysis

which parameters can be quantified, which not where to trigger, what to sample

### **Experimental data**

#### **KINETICS OF SUSTAINED ORAL ADMINISTRATION**

fast rate initial phase
slow rate terminal phase
interindividual variability

NOTE:

sampling scheme
experimental duration
pooled eggs





### **Model reduction**

#### **PBPK modelling approach: from 5 to 2 compartments**

•only sparse and outdated data on body composition

 no data on cardiac output and regional blood flow

 no data on lipid content and lipid composition of tissues (tissue:blood partition coefficients)

•compound property directed reduction: fat compartment for lipophilic compounds





### **Data fitting**

Eggs: 
$$C_{y,f} = C_{y,ss} \left( 1 - \left( a \times e^{\lambda_1 t} + \left( 1 - a \right) \times e^{\lambda_2 t} \right) \longrightarrow$$

Fat:  $C_f = C_{f,ss} (1 - (b \times e^{\lambda_1 t} + (1 - b) \times e^{\lambda_2 t}))$ 



$$\lambda_{1} = -(q_{c} + q_{f} + R + \frac{1}{\sqrt{((q_{c} + q_{f} + R)^{2} - 4q_{f} R))}} / 2$$

$$\lambda_{2} = -(q_{c} + q_{f} + R - \frac{1}{\sqrt{((q_{c} + q_{f} + R)^{2} - 4q_{f} R)}} ) / 2$$

$$a = \frac{\lambda_{2} + R}{\lambda_{1} - \lambda_{2}} \qquad b = \frac{\lambda_{2}}{\lambda_{1} - \lambda_{2}}$$

$$C_{y,ss} = \frac{Y}{W_{y,f}} \frac{F D_{0}}{R} \qquad C_{f,ss} = \frac{q_{c}}{q_{f}} \frac{1}{W_{f}} \frac{F D_{0}}{R}$$

### **Parameter identification**







#### CONDITIONAL





### **Parameter fitting**

#### **TRICK:**

set K = 0, fit parameters:  $Y_{max}$ ,  $F_{min}$ ,  $W_{f,min}$ calculate, assuming  $F_{max} = 1$ ,  $Y_{min}$ ,  $K_{max}$ ,  $W_{f,max}$ validate: setting F = 1, fit  $Y_{min}$ ,  $K_{max}$ ,  $W_{f,max}$ 

#### **RESULT:**

Y: [ 0.043, 0.055 ] day<sup>-1</sup> K: [ 0, 0.011 ] day<sup>-1</sup> F: [ 0.78, 1 ]  $W_f$ : [ 0.23, 0.29 ] kg CALCULATE = VALIDATE

 $F_{max} = 1$ 

# **Modelling result**



### Validation:

 different subgroups : dioxins, furans, mono-ortho PCBs, nonortho PCBs

•10 fold higher exposure & different dioxins composition

non-dioxin like PCBs

### **Application:**

EU-limits in eggs (3 pg TEQ / g yolk fat) and in feed (0.75 ng TEQ / kg feed) do not comply: should be 0.2 ng TEQ / kg feed



### Conclusions

•Modelling transfer of dioxins from feed to eggs succeeded thanks to:

careful experimental set-up

justified choice of kind of model

parameter identification analysis

The model was successfully
 verified on other data
 applied to dioxin limits comparison in eggs and feed





PBPK heaven:  $Q_{eff}$   $CL_h$   $P_c$   $P_{y,f} = P_f$ 

powerful assumption on one parameter value, e.g.  $P_f = 150$ 

or: experimentally determination of one parameter value, e.g.  $P_f$ 

earth:  $q_c q_f Y K$ 

### REDEMPTION

J.C.H. van Eijkeren, M.J. Zeilmaker, C.A. Kan, W.A. Traag & L.A.P. Hoogenboom; A toxicokinetic model for the carry-over of dioxins and PCBs from feed and soil to eggs; *Food Additives and Contaminants*; **23** (2006) 509-517

L.A.P. Hoogenboom, C.A. Kan, M.J. Zeilmaker, J.C.H. van Eijkeren, & W.A. Traag; Carry-over of dioxins and PCBs from feed and soil to eggs at low contamination levels – Influence of mycotoxin binders on the carry over from feed to eggs; *Food Additives and Contaminants*; **23** (2006) 518-527

