

K-PD models as a flexible modeling tool in non-clinical statistics

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Ying, suffering from allergies



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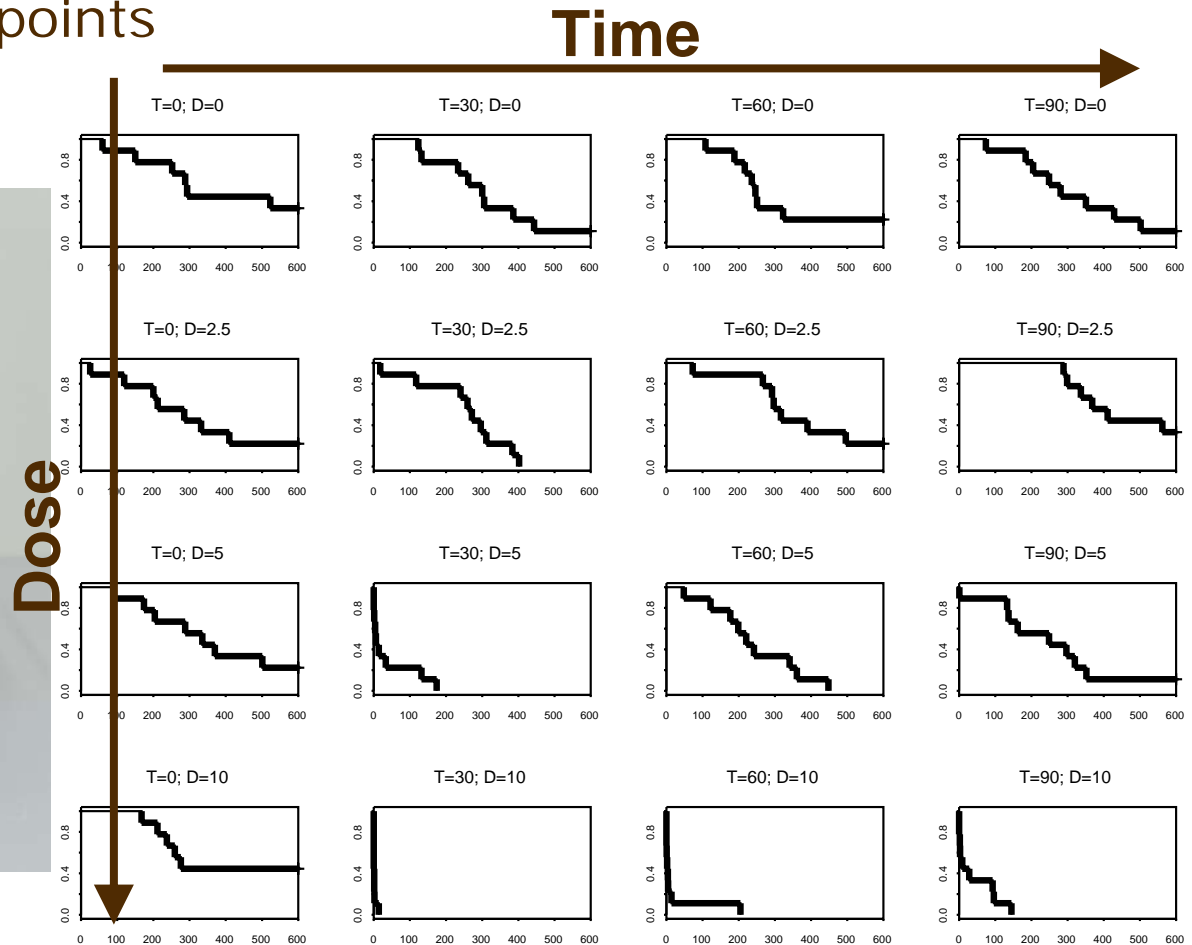
Outline

- Introduction
- Methodology
- Four examples
- Conclusions

Introduction

Pharmacodynamical test: Rotarod (motor coordination)

10 mice/dose, 4 time points



➤ Main question:

How do we quantify the potential effect on motor coordination in a single value?

How do we compare different compounds?

➤ Time to event data

- Parametric: Accelerated Failure Time (AFT), Polynomials
- Nonparametric: Cox regression, AFT with Splines

Methodology

➤ Traditional statistical techniques (e.g. GLM):

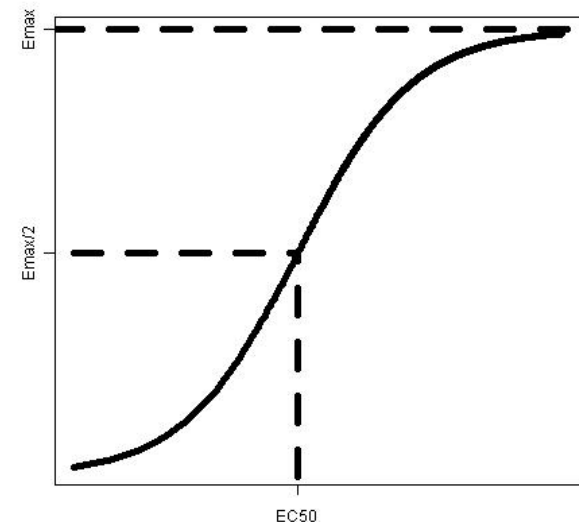
- Polynomials, random effects

$$E(g(Y_{ij}) | b_i, x_{ij}) = \sum_{ij} x'_{ij}\beta + z'_{ij}b_i$$

➤ Quantification of the effect and comparison of compounds?

➤ Alternative approach: PK/PD models

$$E(g(Y_{ij}) | b_i, x_{ij}) = E_{0i} + \frac{E_{max}c_{ij}}{EC_{50} + c_{ij}}$$



KPD

- Fit an Emax model on the scale parameter of the Weibull

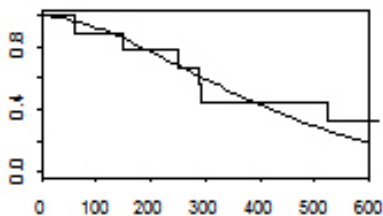
$$E(g(Y_{ij}(t)) \mid b_i, x_{ij}) = \exp(\beta_0) \left[1 - \frac{C_{ij}}{EC_{50} + C_{ij}} \right]$$

- Assume a latent PK-profile

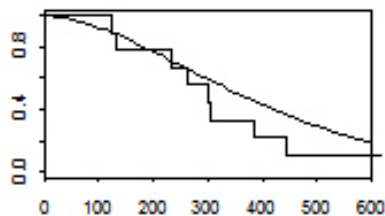
$$C_{ij}(t) = \sum_k A_{ik} \exp(-\theta_{ik}t)$$

- Basically, you assume that an unobserved concentration-time profile at the site of action is driving the response
- Number of compartments depends on the model fit, not on external information or physiology
- Confounding of EC_{50} and C_{ij}

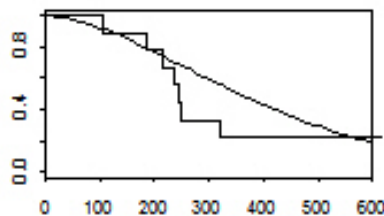
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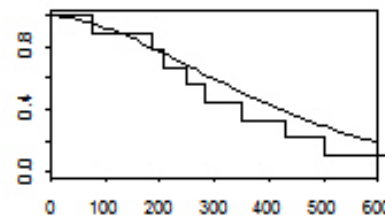
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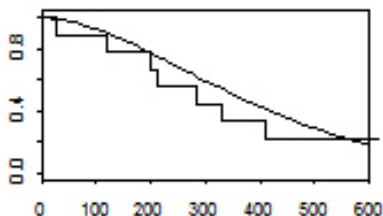
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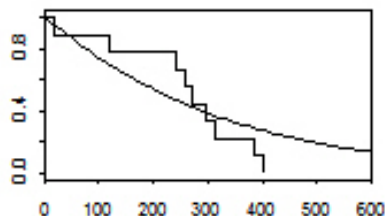
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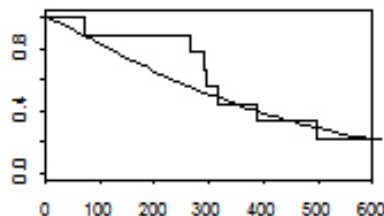
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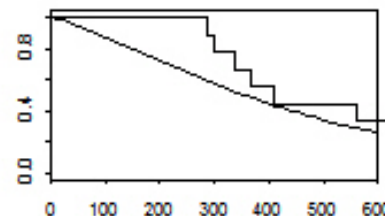
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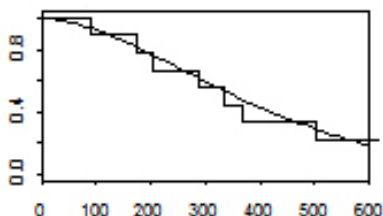
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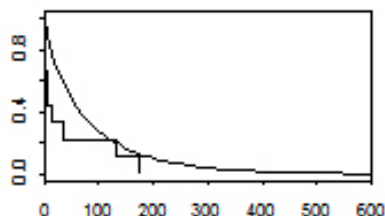
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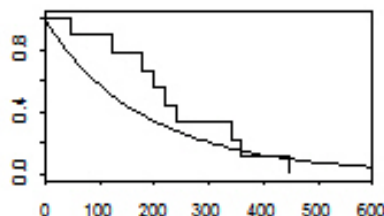
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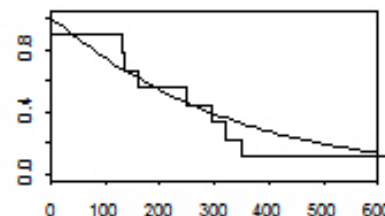
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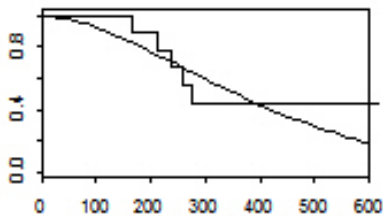
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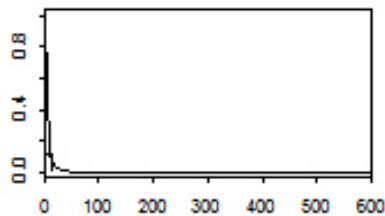
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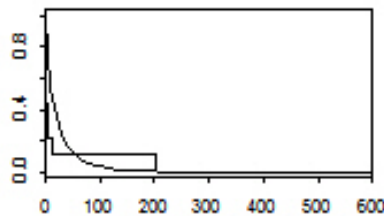
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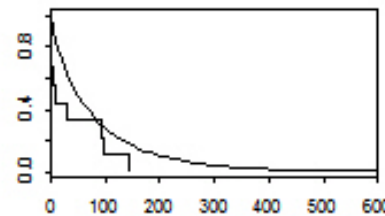
T=30; D=10



T=60; D=10



T=90; D=10



Disease Modeling

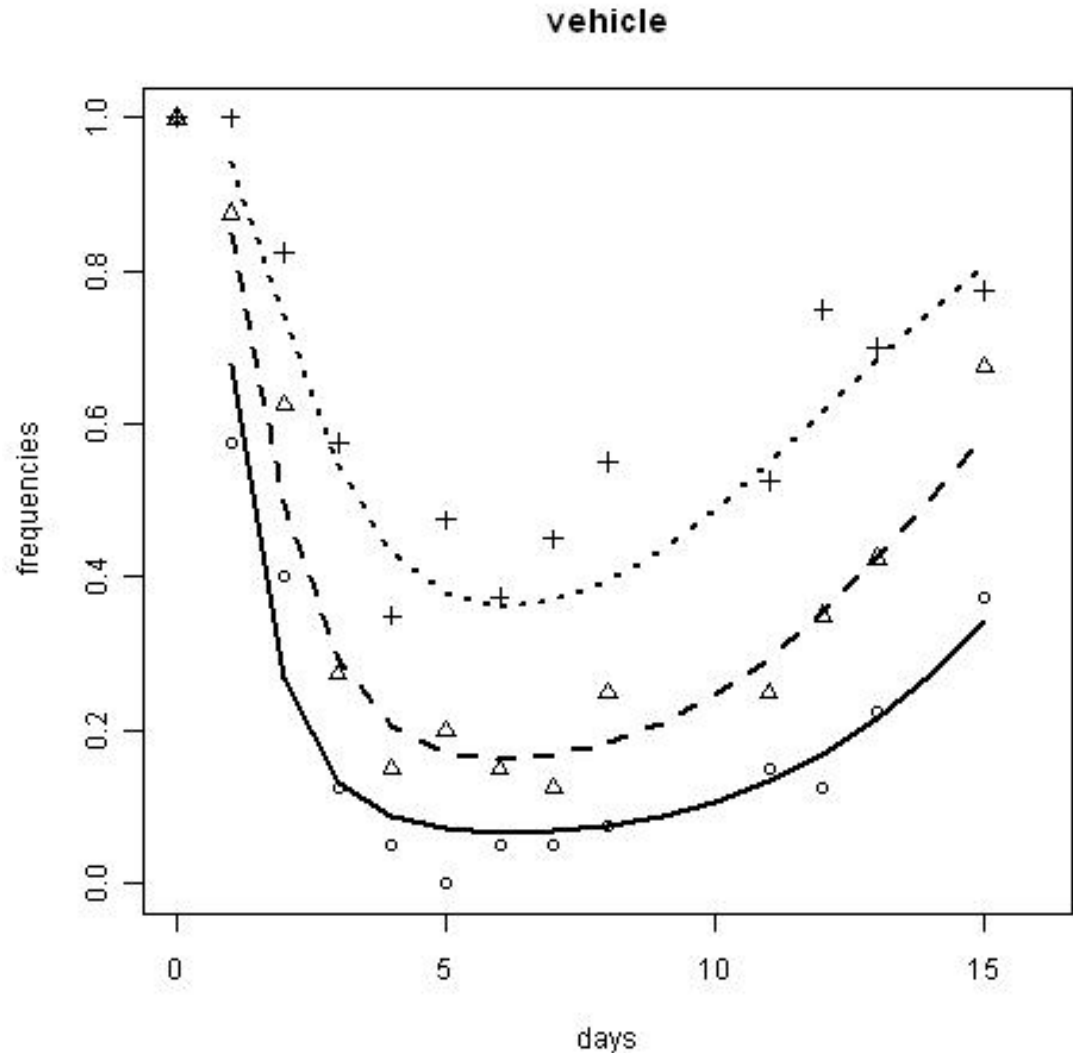
- ▶ What happens if there is a natural evolution over time?
- ▶ Holford states that for clinical data

$$\begin{aligned} & \text{Disease progression} \\ & + \text{Placebo Effect} \longrightarrow E_{0i} = f(t_{ij}, x_{ij}) \\ \frac{E_{max}c_{ij}}{EC_{50} + c_{ij}} \longleftarrow & \text{+ Treatment Effect} \\ & = \text{Response} \longrightarrow E(g(Y_{ij}) | b_i, x_{ij}) \end{aligned}$$

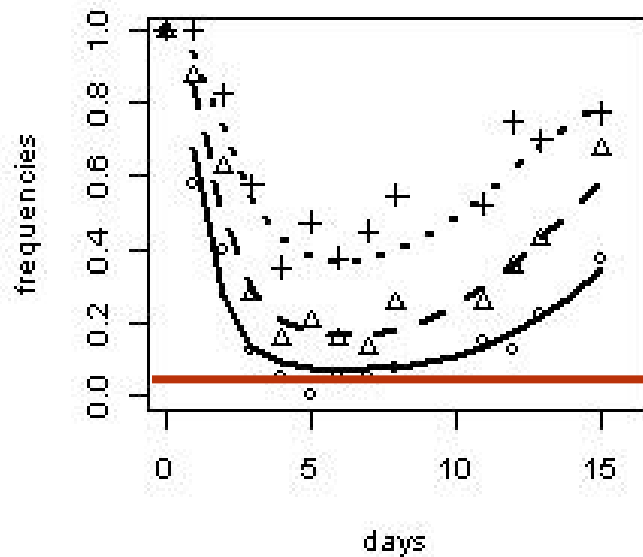
- ▶ Model the vehicle data first, then include the treatment impact
- ▶ Treatment effect can be additive, or included in the parameters of the vehicle model

Pharmacodynamical test: Pin Prick

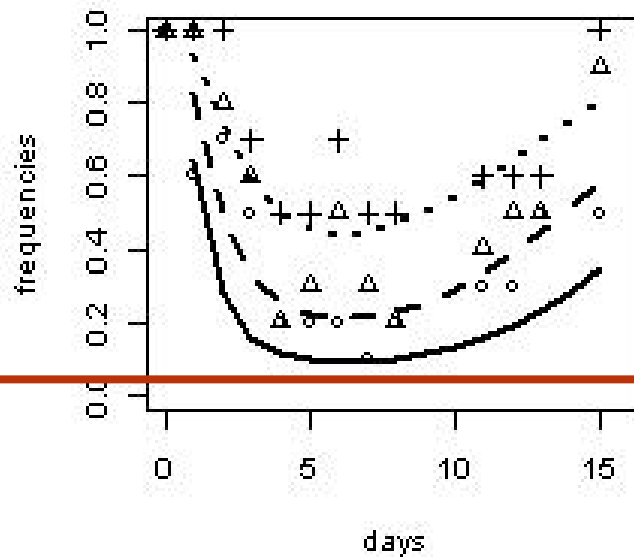
- ▶ 3+1 compounds
add-on to Taxol
- ▶ 10 rats / compound
- ▶ Score: 0, 1, 2, 3
- ▶ How to compare
compounds?
- ▶ Cum-logit prop odds
model (Agresti 2002)
- ▶ Model the impact of
the compounds on
Taxol (E_{max} and/or
 EC_{50})



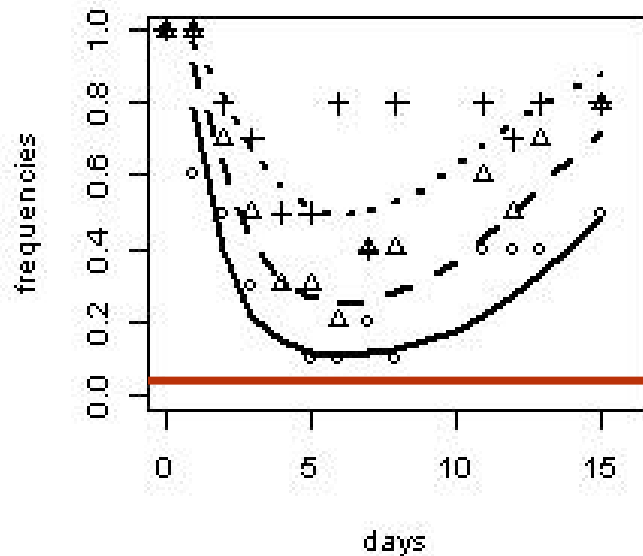
vehicle



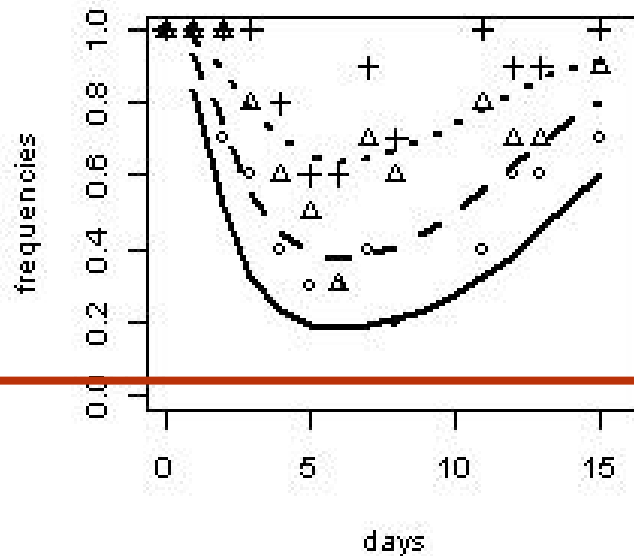
compound 1



compound 2

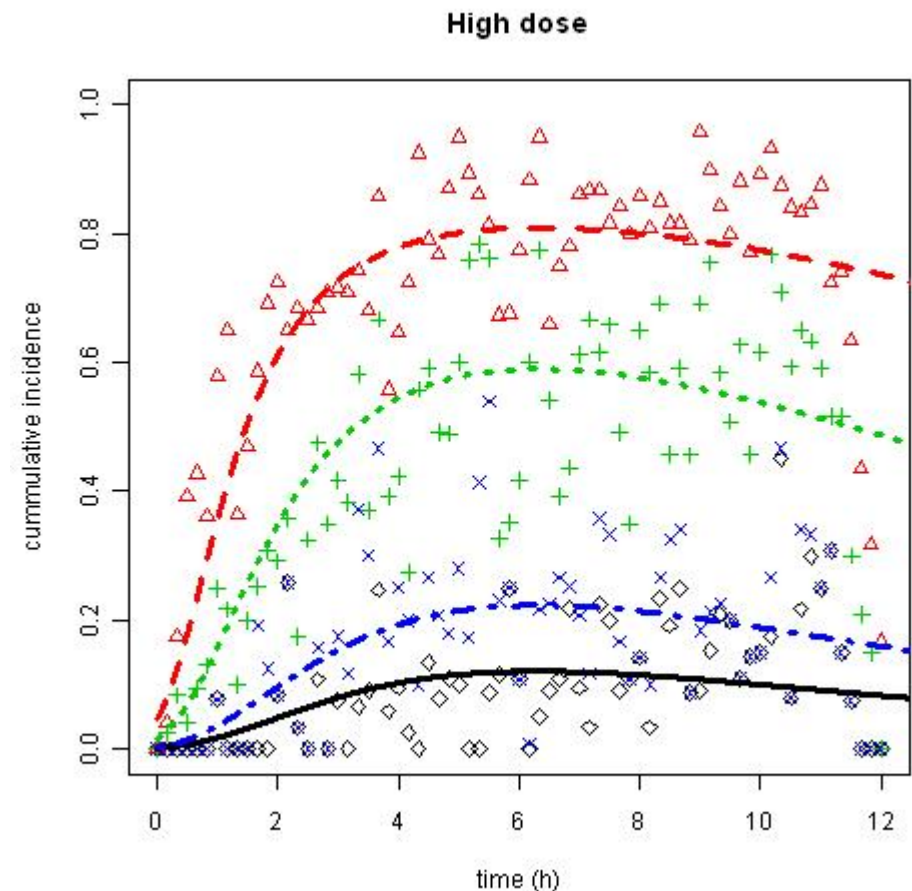
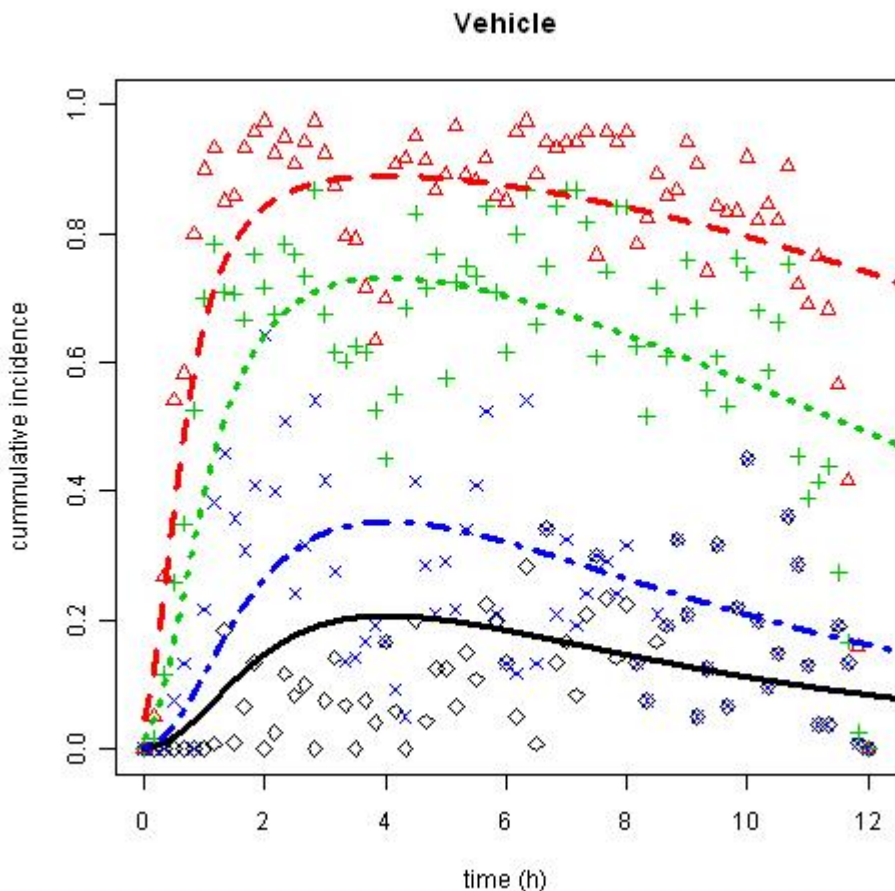


compound 3



Another Pharmacodynamic test

- ⊕ Design
 - 6 animals
 - 4x4 cross-over
 - 12h evaluation
 - Dose response
- ⊕ Test procedure: confidential
- ⊕ Assume a virtual treatment (E_{max} =disease progression + placebo)
- ⊕ Incorporate treatment effect at E_{max} , EC_{50} and PK-parameters
- ⊕ K-PD model for disease progression
- ⊕ Dose as covariate on the absorption coefficient
- ⊕ Quantification of change of onset of effect

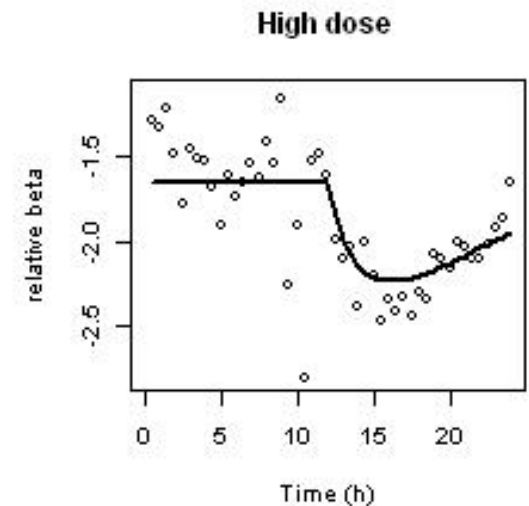
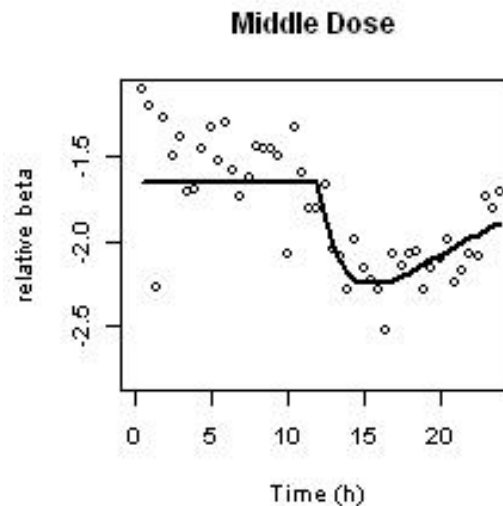
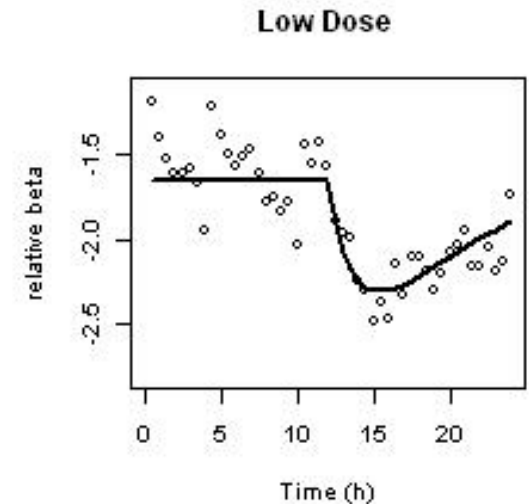
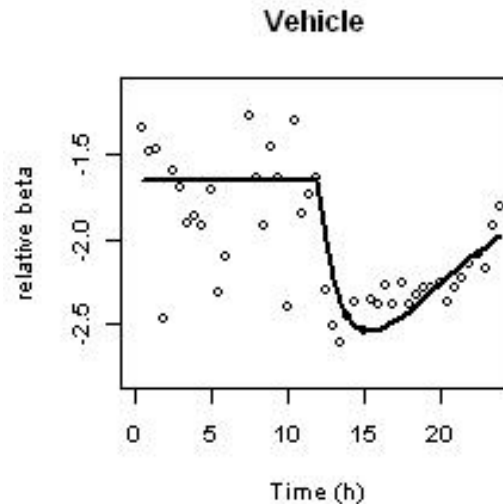


➤ Design

- 6 animals
- 4x4 cross-over
- 24h evaluation
- Dose response

➤ Assume after 12h a virtual treatment (E_{max}=disease progression + placebo)

➤ Incorporate treatment effect at E_{max} and EC₅₀



Are we statisticians? Are we PK-scientists? We are pharmacometricians

- Nonlinear mixed effects modeling has been ignored too long by statisticians
- Interpretation of biologically significant effect
- Flexible modeling allows for model-based drug development
- Enabling optimization and integration of data analysis
- Emphasize **Generalized Nonlinear Mixed Effects**

References

- ▶ Jacqmin P, Snoeck E, van Schaick EA, Gieschke R, Pillai P, Steimer JL, Girard P. *Modelling response time profiles in the absence of drug concentrations: definition and performance evaluation of the K-PD model.* J PK PD. 2007, 34, 57-85.
- ▶ Jacobs T, Straetemans R, Molenberghs G, Bouwknecht JA, Bijmens L. *A Latent Pharmacokinetic Time Profile to Model Dose-Response Survival Data.* J Biopharm Stat, 2010, 20, 759–767.