

# Getting more out of your (eco)toxicity data by applying mechanistic interpretation with open source models



## Toxicokinetic-toxicodynamic modelling – examples to use and available tools

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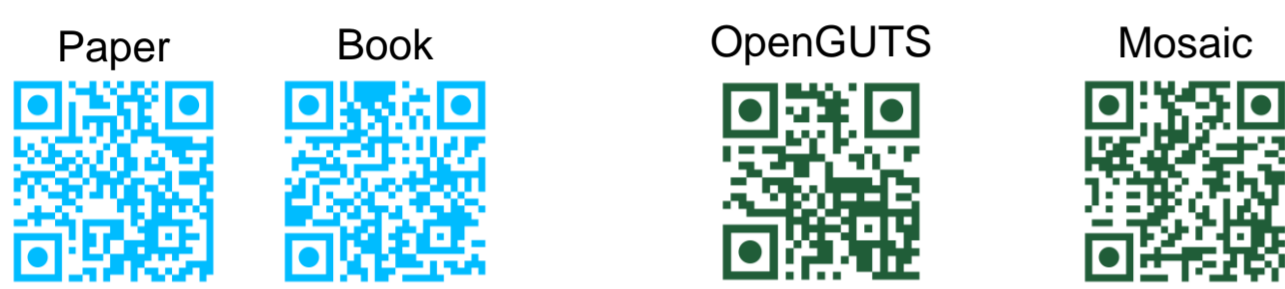
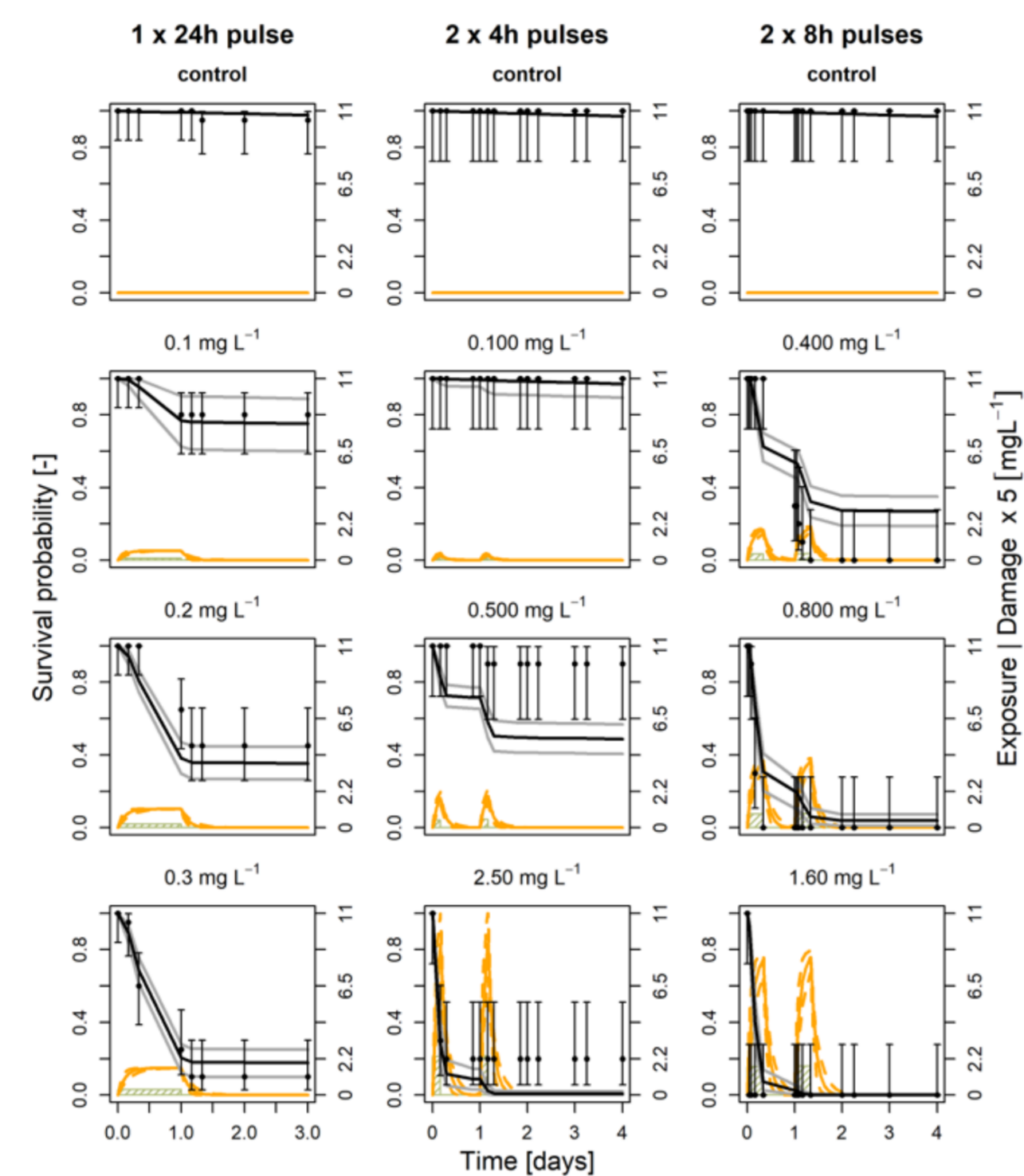
### GUTS

Simulating survival over time at constant or variable exposure.

Species: generic (all species)

Data need: survival over time

→ GUTS model analyse and predict survival under constant and time variable exposure using the whole data package resulting in higher statistical and predictive power

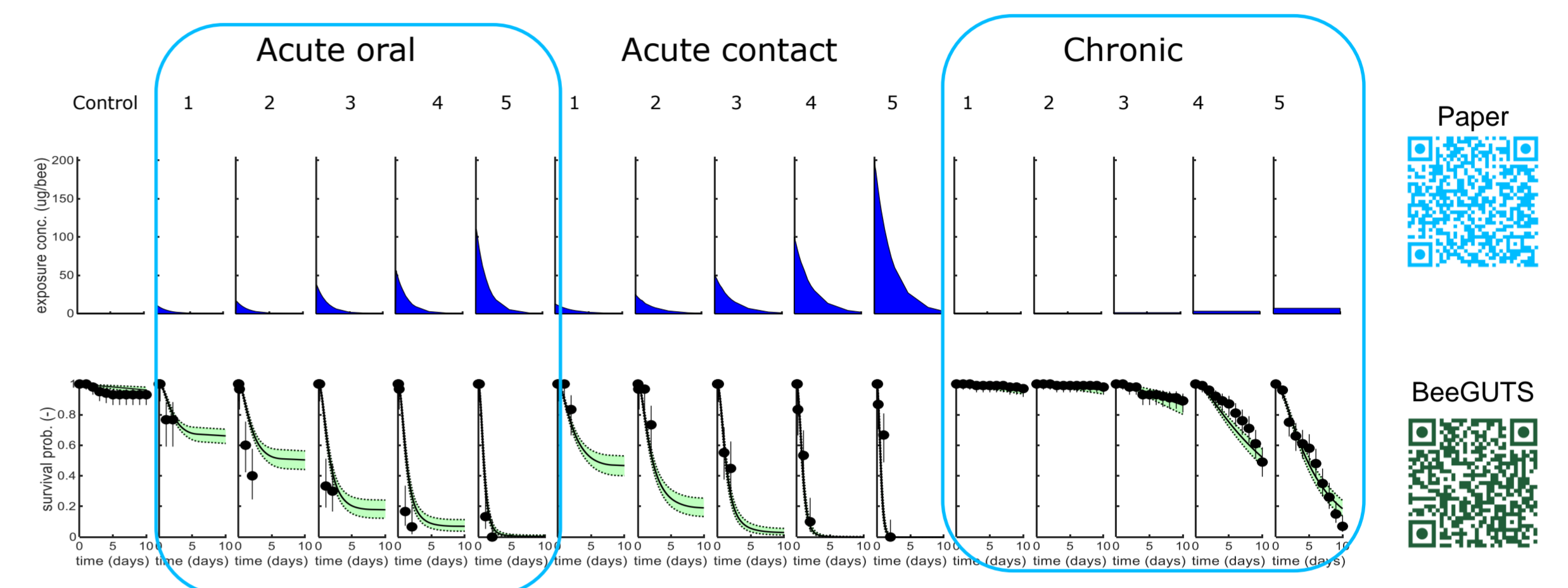


### BeeGUTS

Simulating survival over time for different exposure routes.

Species: honey bee, bumble bee, solitary bees

Data need: survival over time



→ With the BeeGUTS model we are able to analyse all laboratory tests together and extrapolate to real world exposure patterns

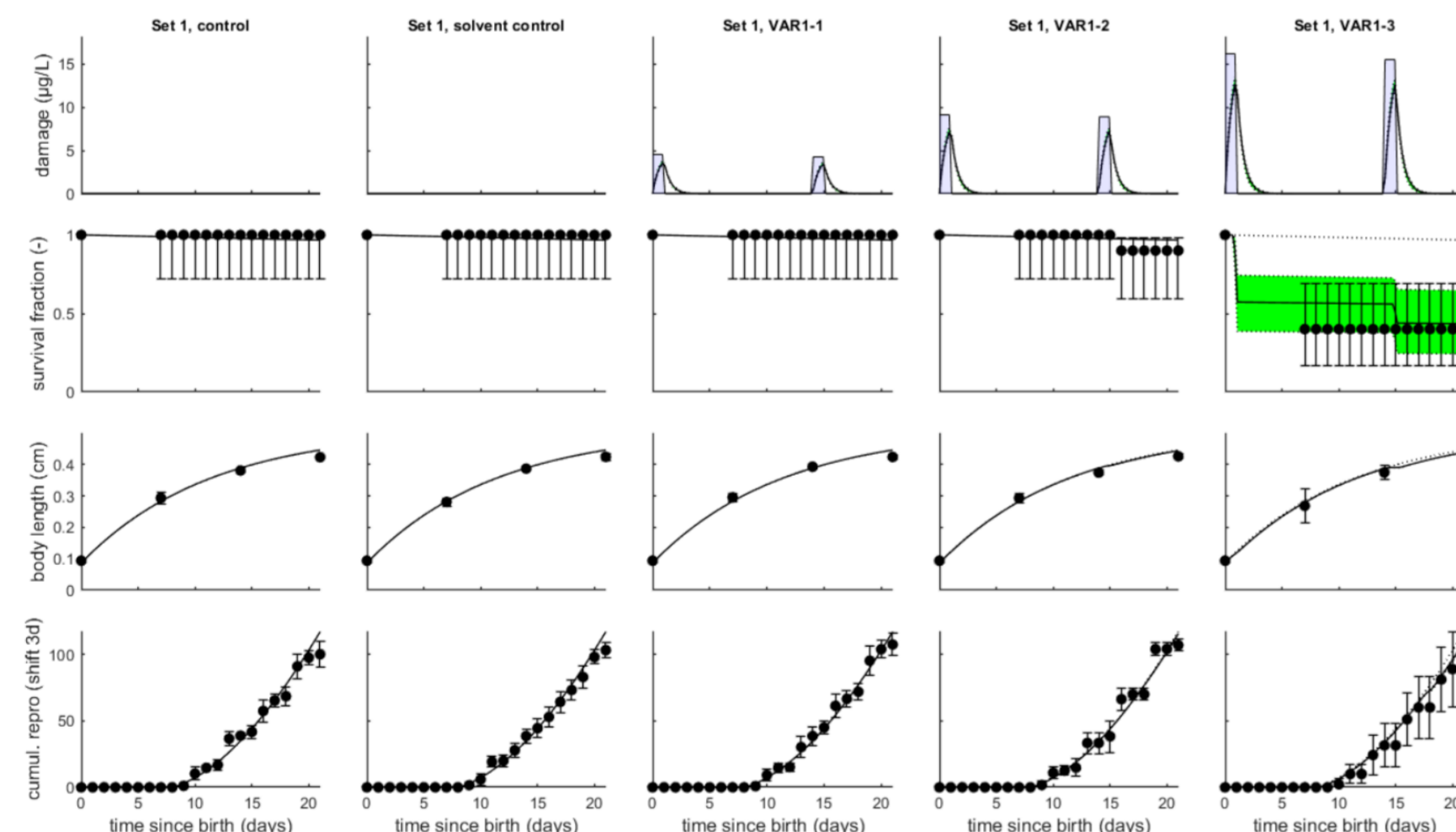


### Dynamic Energy Budget (DEB)

Simulating sublethal effect over time

Species: generic (all animal species)

Data need: effects on reproduction, growth and development



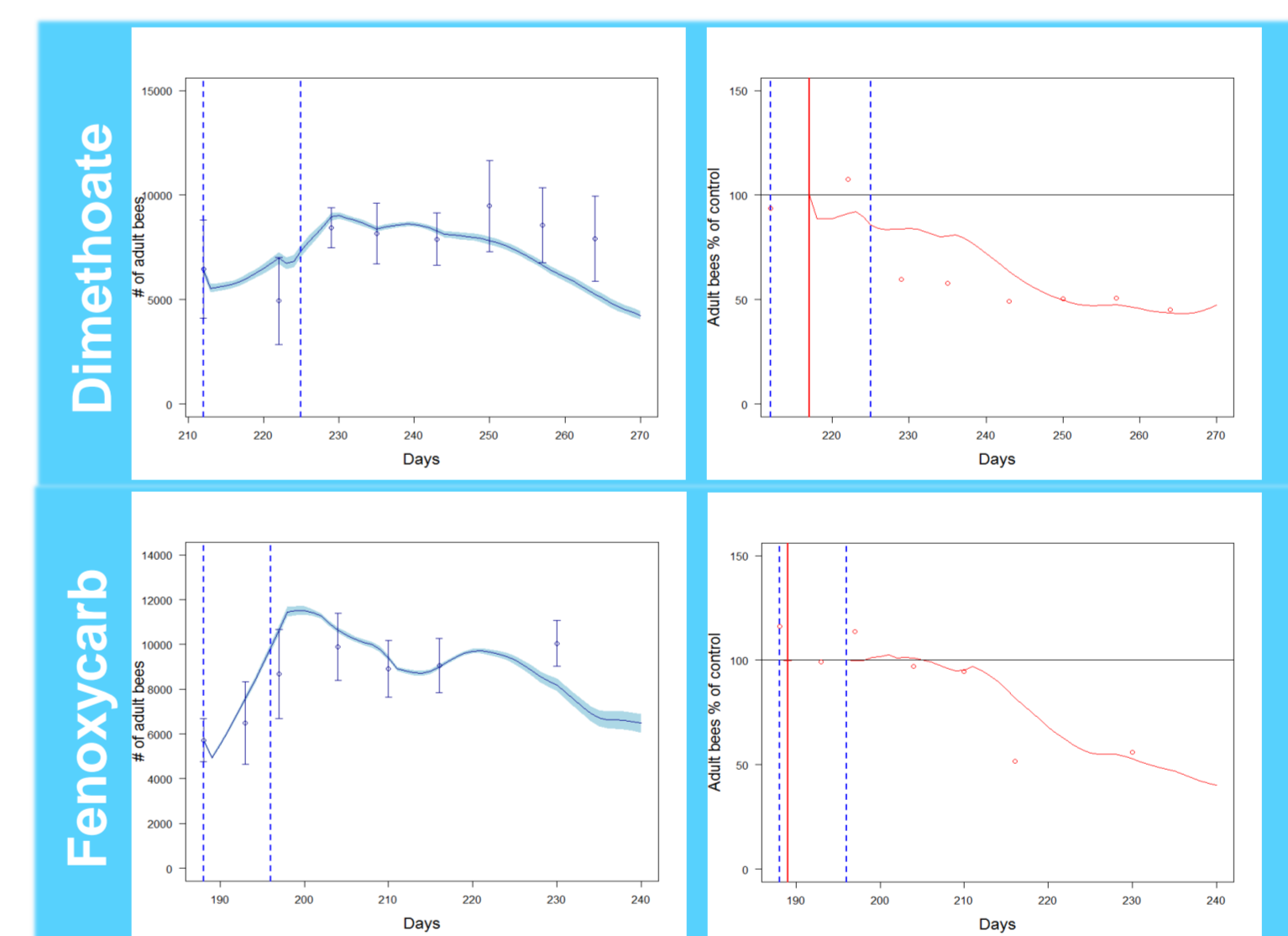
→ With the DEB model effects are analysed in combination and extrapolation to untested exposure scenarios is possible

### BEEHAVE

Simulating colony over time

Species: honey bee

Data need: standard ecotox dataset (OECD 214, 213, 245, 239)



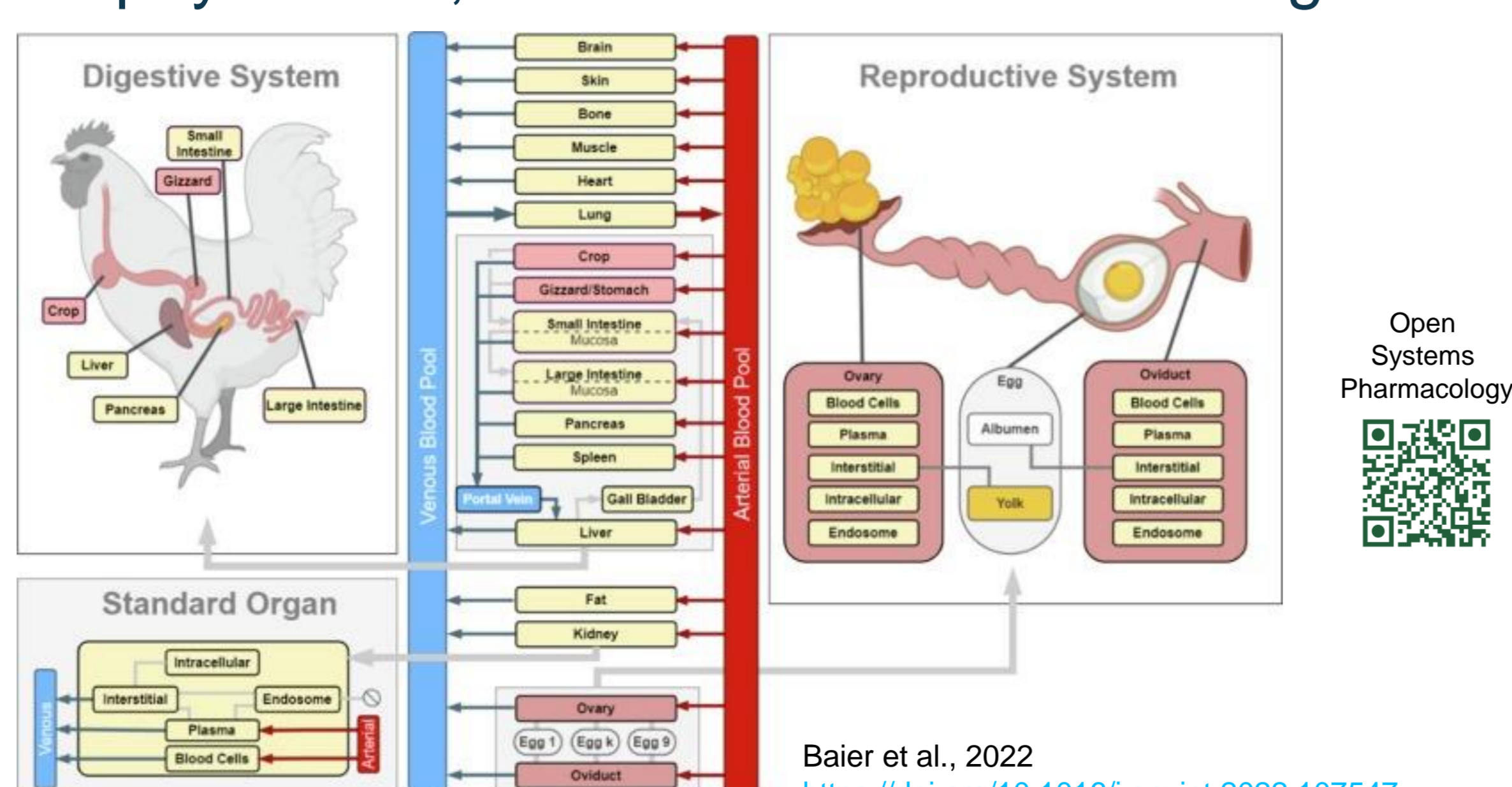
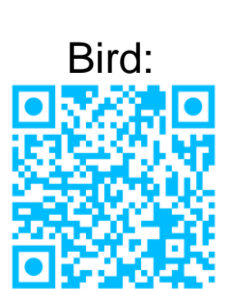
→ Extrapolation from individual (lab) to colony at (semi)field

### PBTK

Simulating uptake, distribution and elimination over time

Species: rat, mouse, dog, human, rabbit, quail, duck

Data need: phys-chem, residues in at least one organ over time



Baier et al., 2022  
<https://doi.org/10.1016/j.envint.2022.107547>  
 created with Biorender

### Why should I use these models?

Standardized TKTD modelling offers a new way of interpreting standard toxicity data. This allows a scientifically sound extrapolation from the constant laboratory condition to the local situation.

This approach allows to extrapolate to different environmental conditions and maybe even to untested species.