

Ready-to-use modelling and statistical tools for advanced environmental risk assessment



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Toxicity tests carried out within laboratories, according to standardized protocols



Appropriate and reliable mathematical models and statistical inference methods



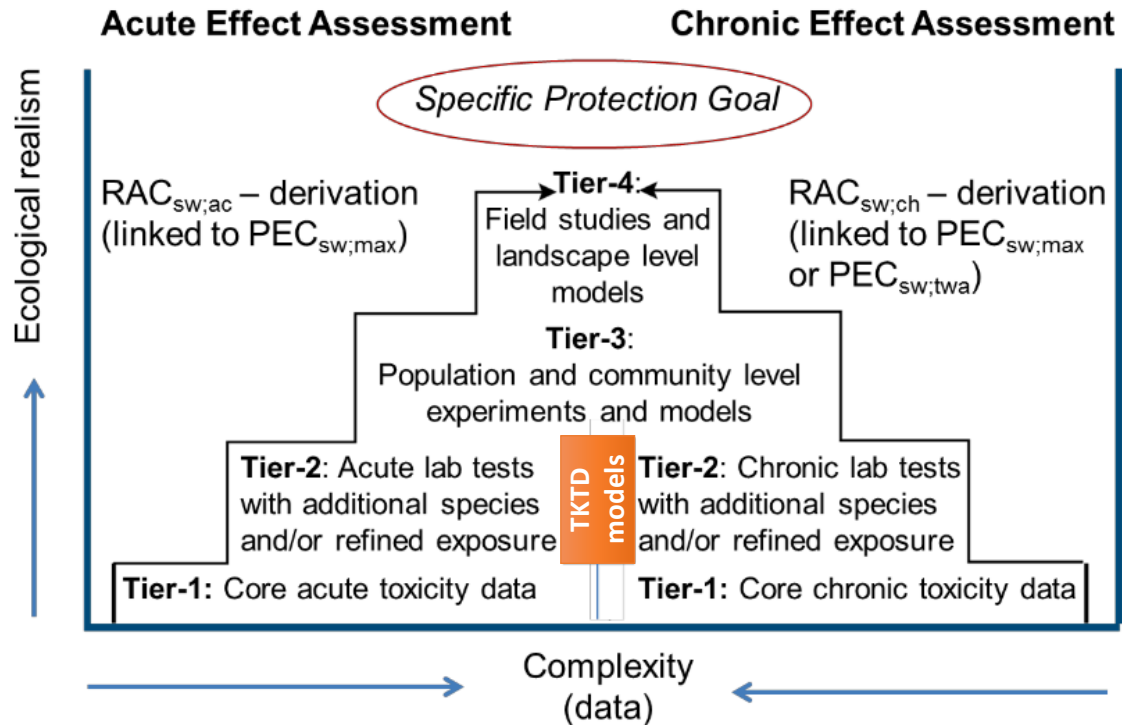
Exposure-response/effect models → Classical outputs such as LC_x/EC_x estimates



Toxicokinetic-toxicodynamic (TKTD) models → advanced analyses accounting for the **time-dependency** of the data.

Environmental Risk Assessment (ERA)

2013 – Aquatic Guidance Document



**Scientific Opinion on the state of the art of
Toxicokinetic/Toxicodynamic (TKTD) effect models for
regulatory risk assessment of pesticides for aquatic
organisms**

EFSA Panel on Plant Protection Products and their Residues (PPR),
Colin Ockleford, Paulien Adriaanse, Philippe Berny, Theodorus Brock, Sabine Duquesne,
Sandro Grilli, Antonio F. Hernandez-Jerez, Susanne Hougaard Bennekou, Michael Klein,
Thomas Kuhl, Ryszard Laskowski, Kyriaki Machera, Olavi Pelkonen, Silvia Pieper,
Robert H. Smith, Michael Stemmer, Ingvar Sundh, Aldrik Tiktak, Christopher J. Topping,
Gerrit Wolterink, Nina Cedergreen, Sandrine Charles, Andreas Focks, Melissa Reed,
Maria Arena, Alessio Ippolito, Harry Byers and Ivana Teodorovic



European Food Safety Authority

Main conclusions

- **GUTS models** (General Unified Threshold models of Survival) for lethal effects of pesticides on animals are well established and can be used in the ERA scheme when exposure varies over time.
- **DEBtox models** (Dynamic Energy Budget for ecotoxicity) for sublethal effects of pesticides on growth and reproduction are considered to be in an advanced state but not yet ready-to-use for ERA.
- Among species-specific models accounting for the effects of pesticides on **primary producers**, the *Lemna* model is suitable for use in ERA while some shortcomings prevent to recommend the *Myriophyllum* and algae models as fit-for-purpose.

1 Calibration

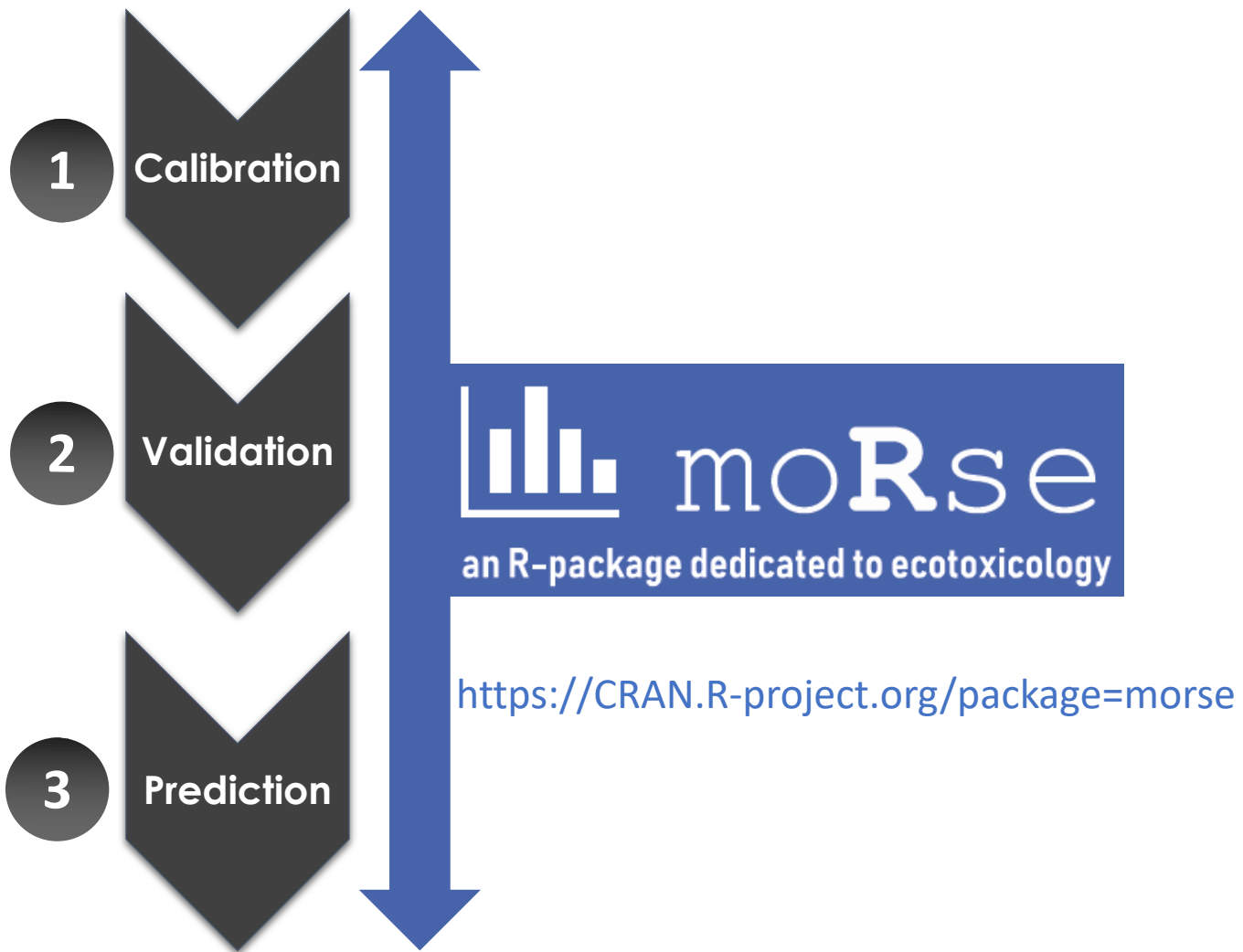
Fit a TKTD model on toxicity test data and get parameter estimates associated with their uncertainty

2 Validation

Simulate an effect over time under a time-variable exposure profile and compare with observed data from a refined toxicity test
→ Three validation criteria are recommended by EFSA

3 Prediction

Make simulations under realistic scenarios to assess risk on how far is the exposure profile from causing a pre-defined effect.
Using GUTS models → Concept of the $x\%$ Lethal Profile (LP x) = Multiplication Factor leading to an additional $x\%$ reduction in the final survival rate.

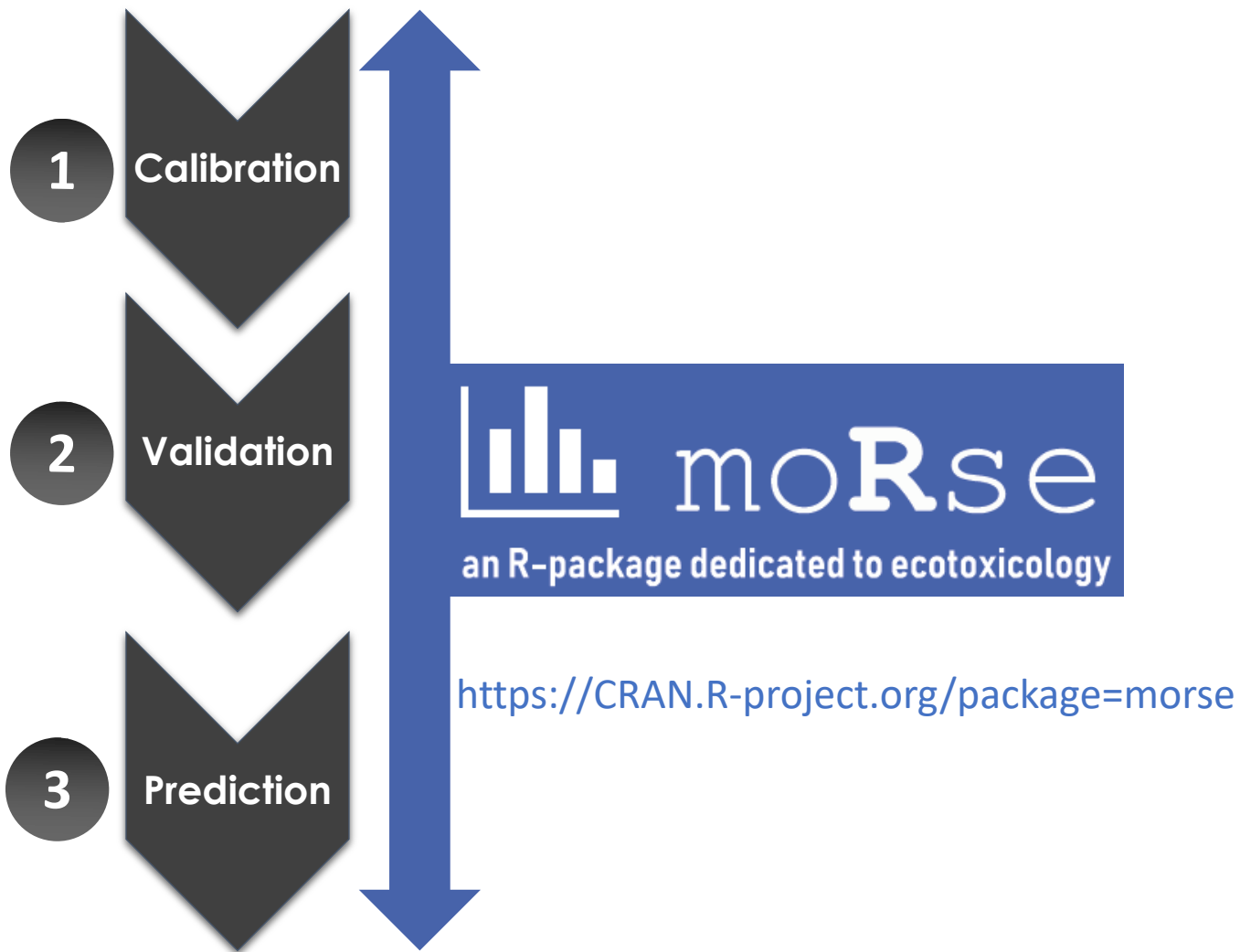


Ready-to-use tools: GUTS as a case-study

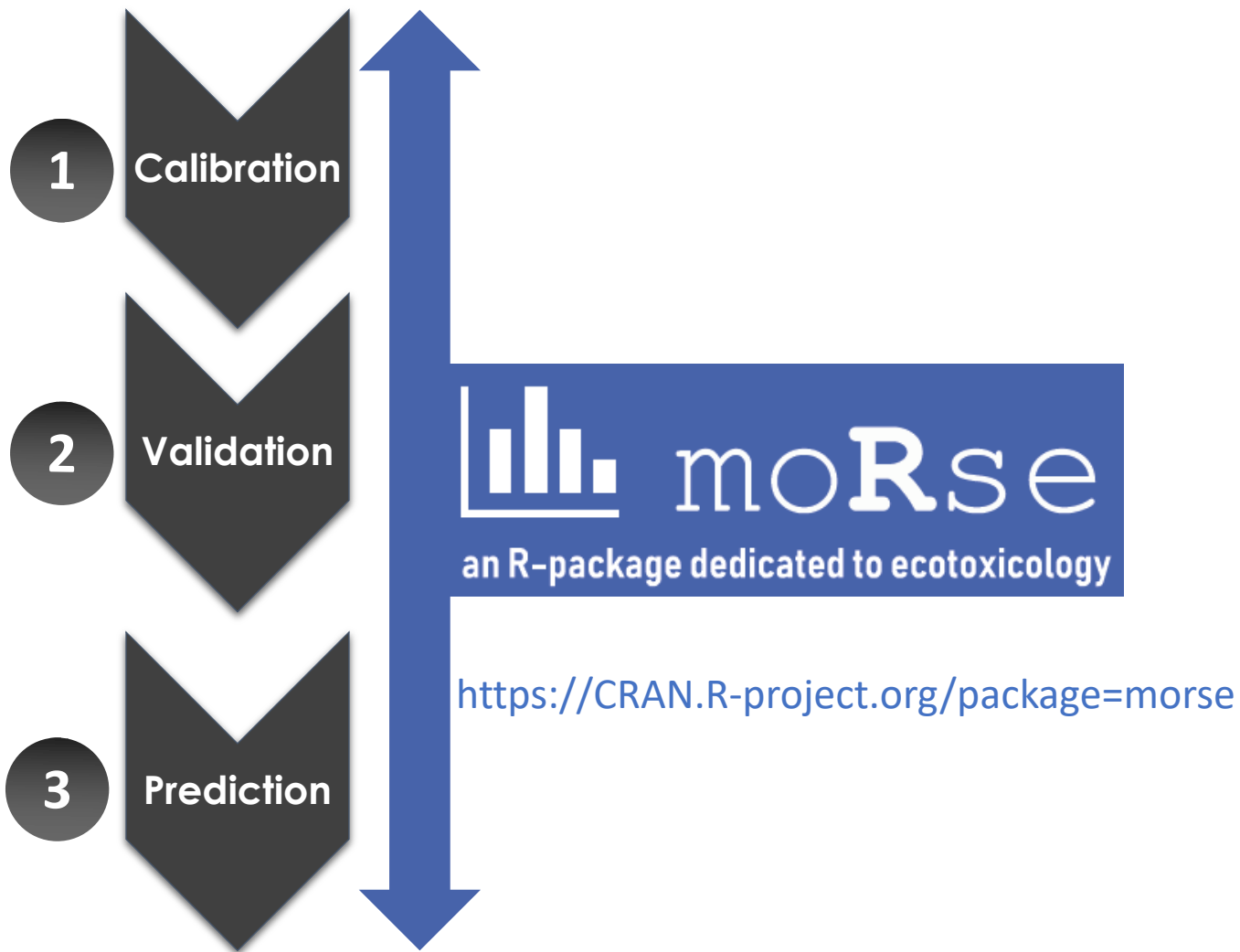
6th International Symposium of DEB theory

March 2019

6



Ready-to-use tools: GUTS as a case-study



Ready-to-use tools: GUTS as a case-study



The R-package moRse

Calibration of classical exposure-response/effect models to get LCx/ECx estimates associated with the quantification of their uncertainty through 95% credible intervals within a **Bayesian framework**;

1. **Calibration** of the General Unified Threshold model of Survival (GUTS) to get parameter estimates for both SD or IT reduced versions, as for example the No Effect Concentration (NEC) or the dominant rate constant (k_d);
2. **Validation** of GUTS model outputs by comparison with observed data, based on results on parameter estimates from a GUTS model calibration;
3. **Prediction** of lethal effects under realistic time-variable exposure concentration scenarios to get x% Lethal Profiles (LPx) estimates, that is the multiplication factor applied to the profile that leads to x% of reduction in the final survival rate.



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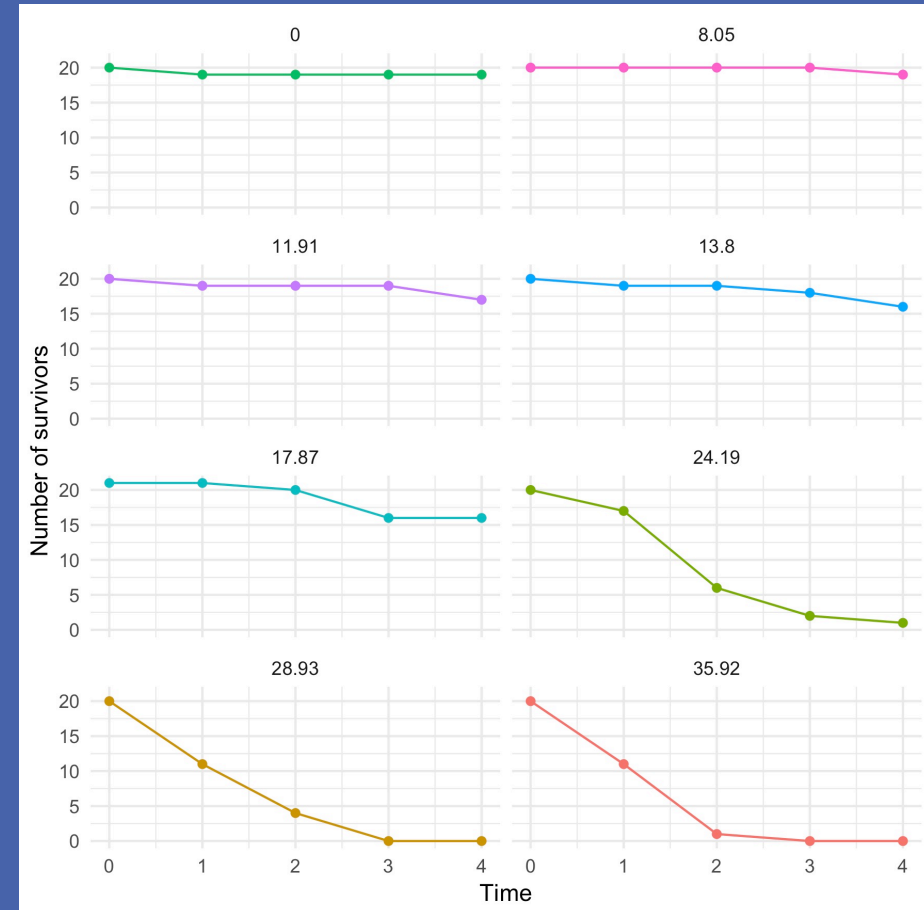
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```
# (1) Load the 'morse' package in your R session  
library('morse')
```

GUTS analyses – 1 Calibration



```
# (2) Load an example (survival data set)
data("propiconazole")
# (3) Create a surv morse object
sdat <- survData(cadmium2)
# (4) Plot raw data (survival)
plot(sdat)
```

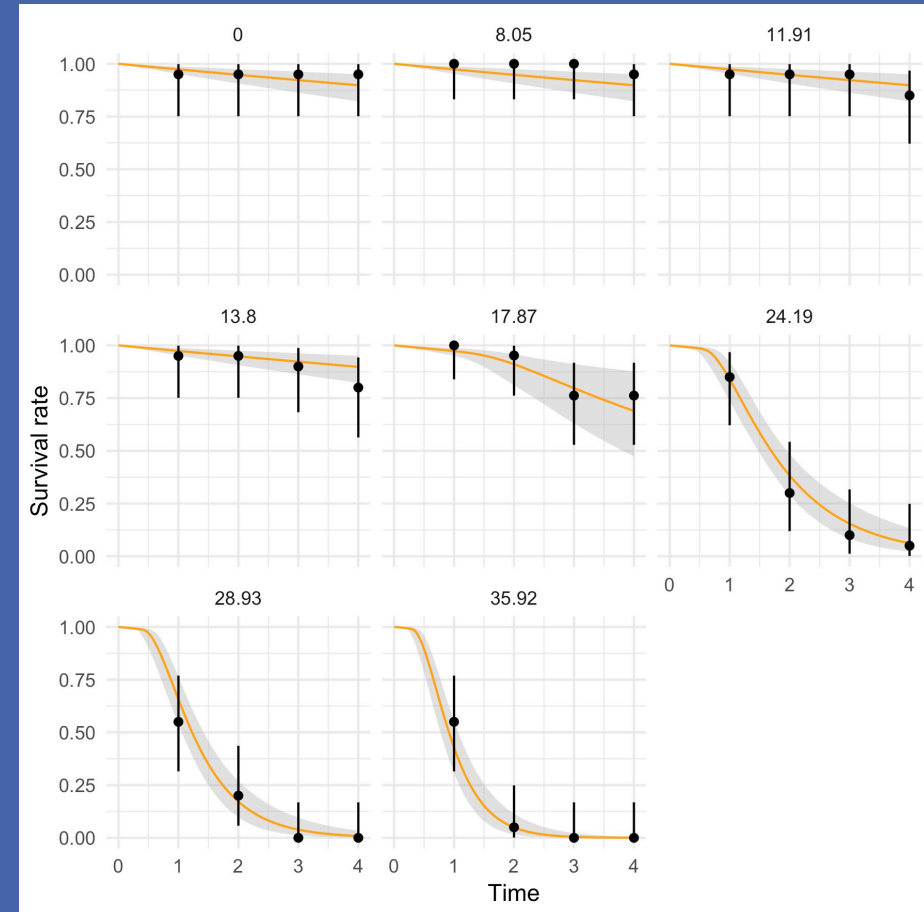


Nyman, A.-M., Schirmer, K., Ashauer, R., (2012) Toxicokinetic-toxicodynamic modelling of survival of *Gammarus pulex* in multiple pulse exposures to propiconazole: model assumptions, calibration data requirements and predictive power. *Ecotoxicology*, (21), 1828-1840.

GUTS analyses – 1 Calibration



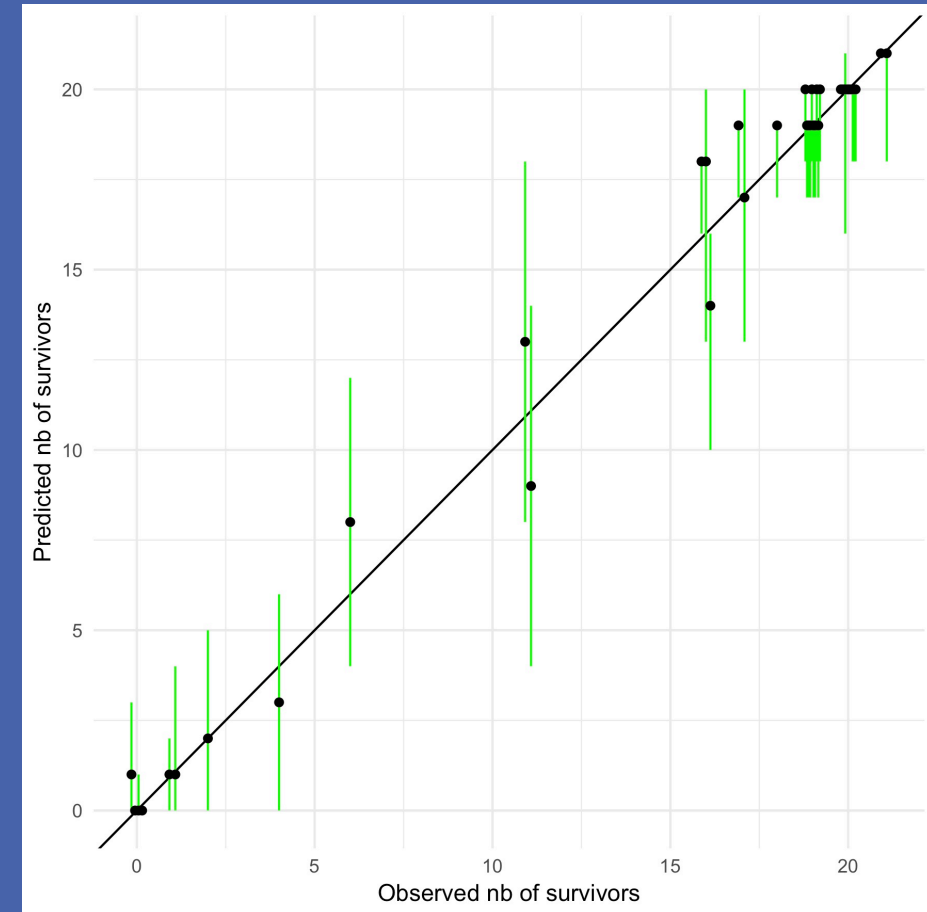
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plot(sdatt)
# (5) Fit a GUTS-RED-SD model
sfit <- survFit(sdatt, model_type = 'SD')
# (6) Get model parameter estimates
summary(sfit)
# (7) Plot fit
plot(sfit, xlab = "Time", adddata = TRUE)
```



GUTS analyses – 1 Calibration



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plot(sfit, xlab = "Time", adddata = TRUE)
# (8) Check goodness-of-fit
ppc(sfit)
```

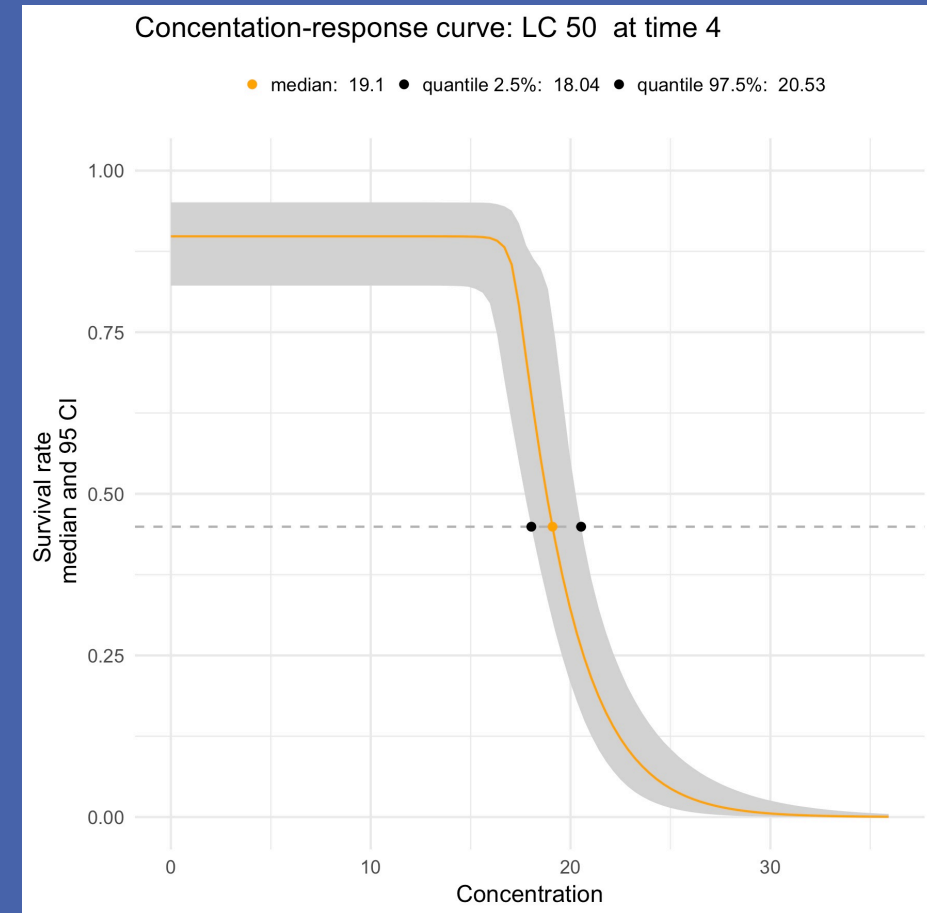


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summary(sfit)
# (7) Plot fit
plot(sfit, xlab = "Time", adddata = TRUE)
# (8) Check goodness-of-fit
ppc(sfit)
# (9) Predict the LC50 at final time
LCx(sfit, X = 50)$df_LCx
plot(LCx(sfit, X = 50))

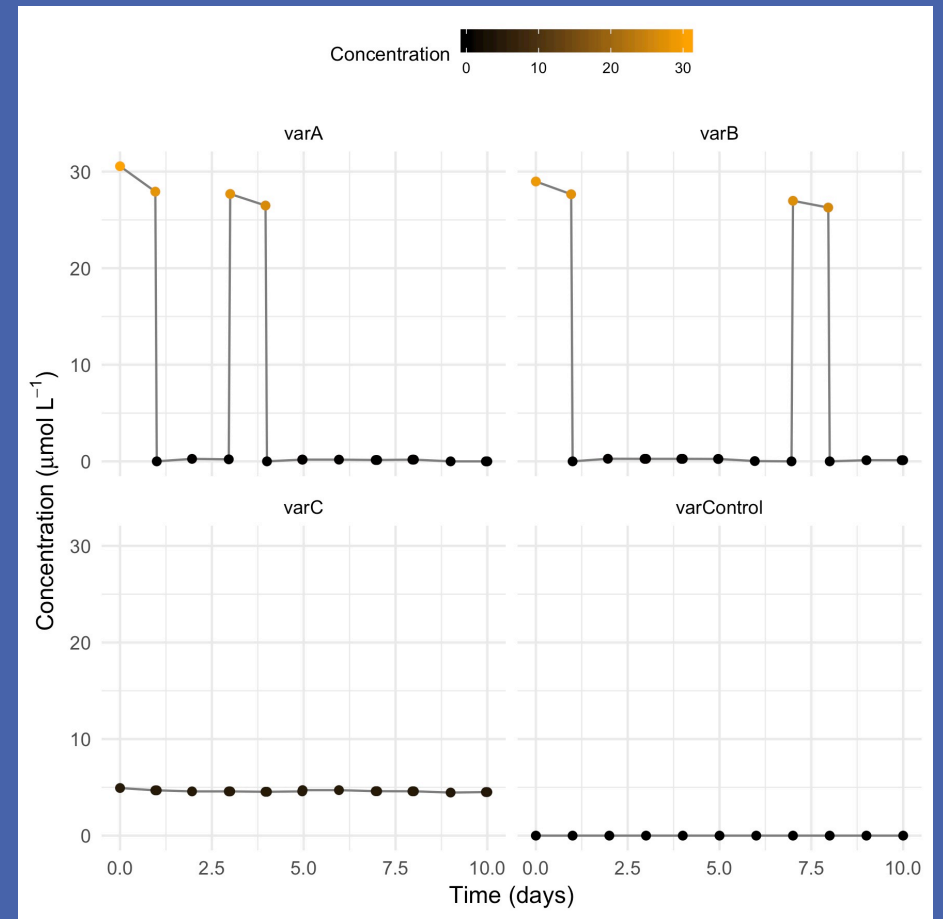
```



GUTS analyses – 2 Validation



```
# (10) Exposure profile  
data("propiconazole_pulse_exposure")
```

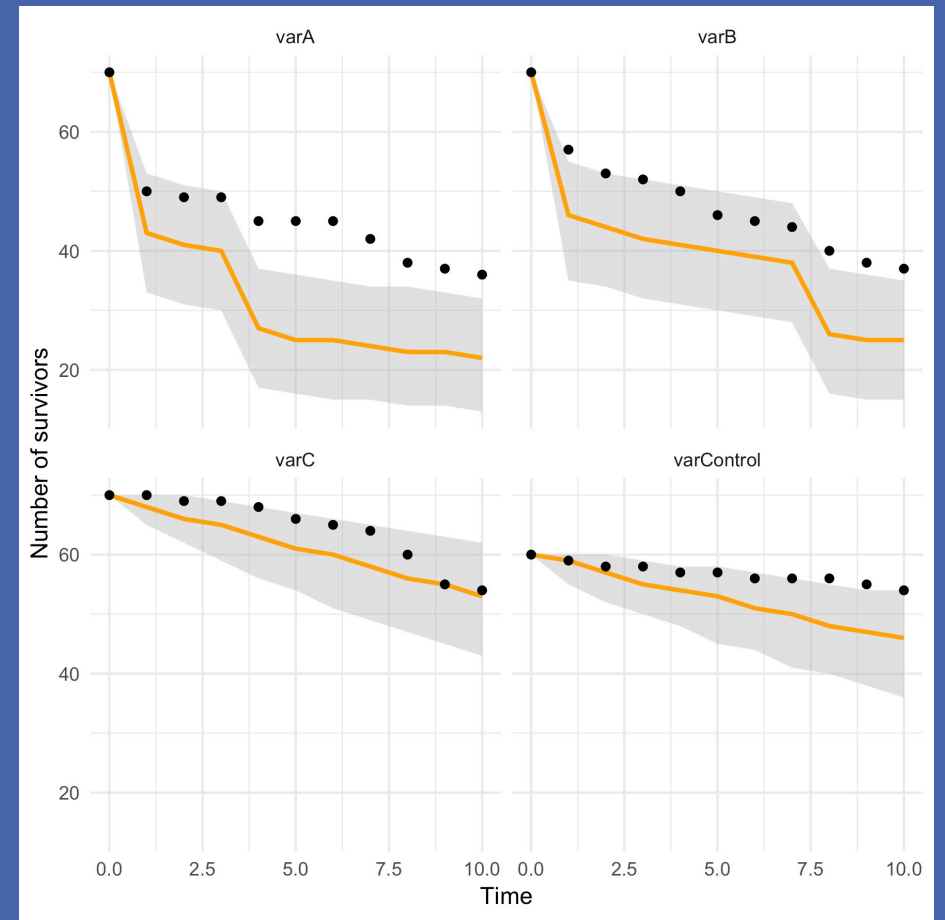


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GUTS analyses – 2 Validation



```
# (10) Exposure profile
data("propiconazole_pulse_exposure")
# (11) Predict the number of survivors
predict_Nsurv_cstT0var <- predict_Nsurv(sfit,
propiconazole_pulse_exposure)
### Note that computing can be quite long
### (until several minutes)
# (12) Plot validation
plot(predict_Nsurv_cstT0var)
```



GUTS analyses – 2 EFSA validation criteria



```
# (10) Exposure profile
data("propiconazole_pulse_exposure")
# (11) Predict the number of survivors
predict_Nsurv_cstTOver <- predict_Nsurv(sfit,
propiconazole_pulse_exposure)
### Note that computing can be quite long
### (until several minutes)
# (12) Plot validation
plot(predict_Nsurv_cstTOver)
# (13) EFSA validation criteria
predict_Nsurv_check(predict_Nsurv_cstTOver)
```

```
$Percent_PPC
  replicate      PPC
1      varA    36.36364
2      varB    63.63636
3      varC   100.00000
4 varControl    81.81818

$Percent_PPC_global
[1] 70.45455
```

Percentage of observations within the predicted uncertainty limits.

Based on experience, results less than 50% of the observations within the uncertainty limits indicate poor model performance.

GUTS analyses – 2 EFSA validation criteria



```
# (10) Exposure profile
data("propiconazole_pulse_exposure")
# (11) Predict the number of survivors
predict_Nsurv_cstTOvar <- predict_Nsurv(sfit,
propiconazole_pulse_exposure)
### Note that computing can be quite long
### (until several minutes)
# (12) Plot validation
plot(predict_Nsurv_cstTOvar)
# (13) EFSA validation criteria
predict_Nsurv_check(predict_Nsurv_cstTOvar)
```

```
$Percent_NRMSE
  replicate      NRMSE
1         varA  31.153292
2         varB  19.959361
3         varC   5.853125
4   varControl   9.130619

$Percent_NRMSE_global
[1] 17.08304
```

Normalized Root Mean Square Error expressed in %.

Based on experience, it is expected that the NRMSE should not exceed the upper limit of 50%.

GUTS analyses – 2 EFSA validation criteria



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# (10) Exposure profile
data("propiconazole_pulse_exposure")
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predict_Nsurv_cstT0var <- predict_Nsurv(sfit,
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### Note that computing can be quite long
### (until several minutes)
# (12) Plot validation
plot(predict_Nsurv_cstT0var)
# (13) EFSA validation criteria
predict_Nsurv_check(predict_Nsurv_cstT0var)
```

\$Percent_SPPE		
	replicate	NRMSE
1	varA	20.000000
2	varB	17.142857
3	varC	1.428571
4	varControl	13.333333

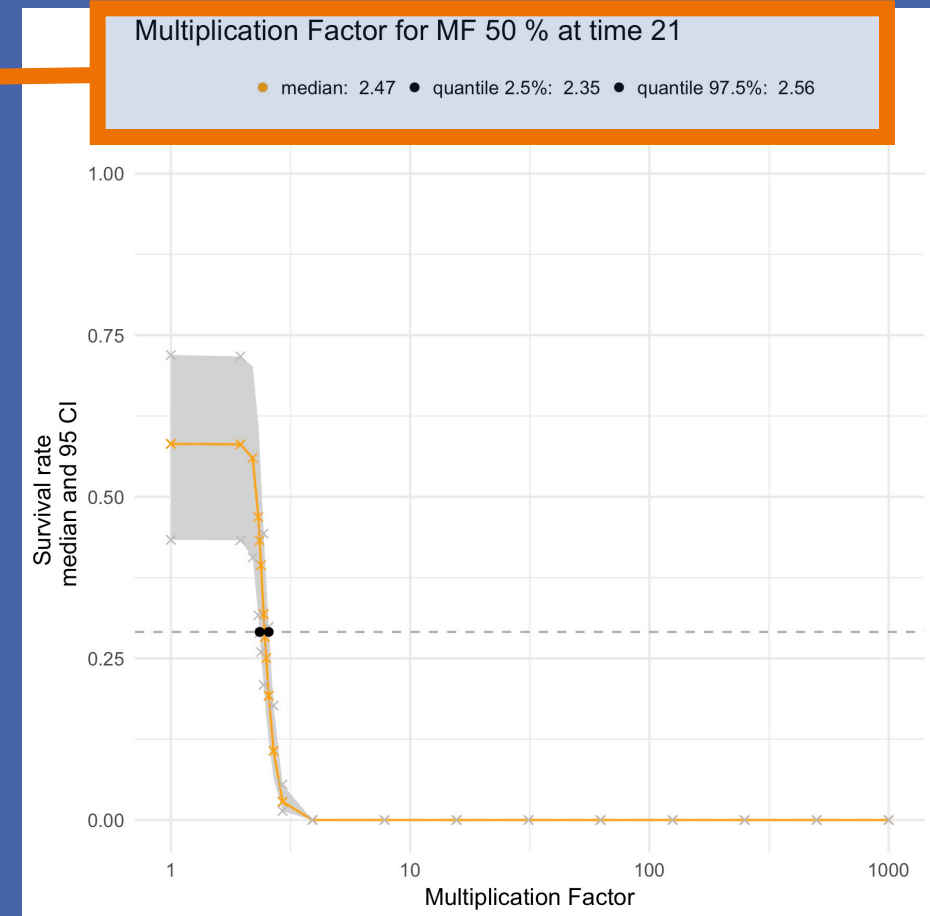
Survival Probability Prediction Error standing for model accuracy considering survival probabilities **at the end** of the exposure profile.

The SPPE value is negative for an underestimation of effects, else it is positive.

GUTS analyses – 3 Predictions

```
# (14) Import the exposure scenario  
# (15) Estimate the MF50 at final time  
MF50 <- MFx(object = sfit, data_predict  
= scenario, quiet = TRUE)  
MF50$df_MFx  
plot(MF50, log_scale = TRUE)
```

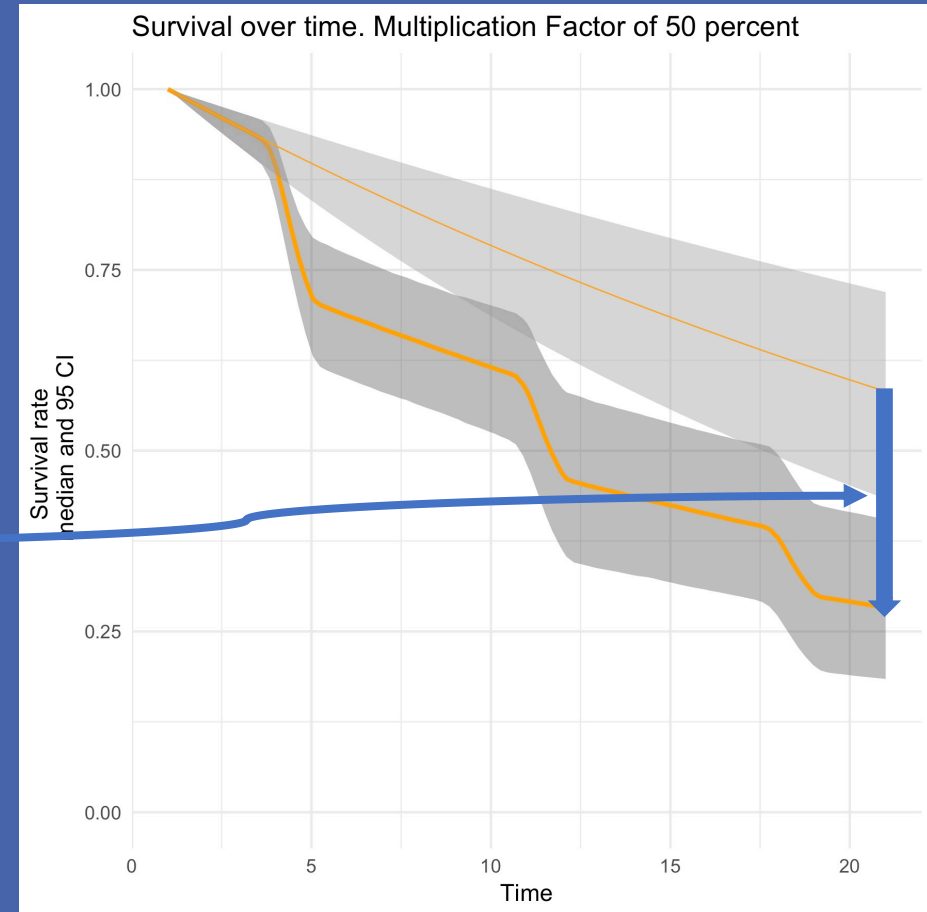
LP₅₀ = 2.47 [2.35 ; 2.56]



GUTS analyses – 3 Predictions

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# (14) Import the exposure scenario  
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MF50 <- MFx(object = sfit, data_predict  
= scenario, quiet = TRUE)  
MF50$df_MFx  
plot(MF50, log_scale = TRUE)  
plot(MF50, x_variable = "Time")
```

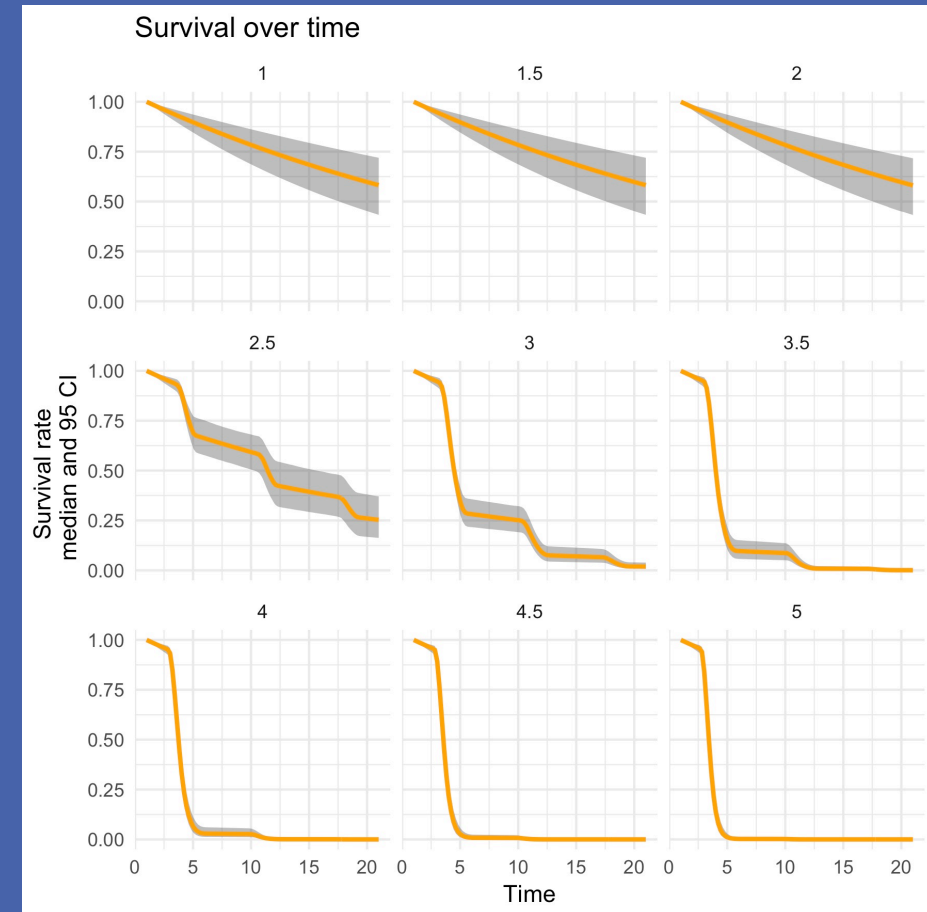
$LP_{50} = 2.47 [2.35 ; 2.56]$



GUTS analyses – 3 Predictions



```
# (14) Import the exposure scenario
# (15) Estimate the MF50 at final time
MF50 <- MFx(object = sfit, data_predict
= scenario, quiet = TRUE)
MF50$df_MF
plot(MF50, log_scale = TRUE)
plot(MF50, x_variable = "Time")
# (16) Plot the MF-response curve
MF <- MFx(object = sfit, data_predict =
scenario, X = NULL, MFx_range =
seq(1, 5, 0.5))
plot(MF)
# (17) Plot the survival rate over time
for various MF
MF <- MFx(object = sfit, data_predict =
scenario, X = NULL, MFx_range =
seq(1, 3.5, 0.5))
plot(MF, x_variable = "Time")
```



**All the previous analyses
can be done directly on-line**



<http://pbil.univ-lyon1.fr/software/mosaic/>

Video-01

**Few seconds/minutes
later...**

EFSA workflow: calibration (step 1)

Video-02

EFSA workflow steps 2 and 3



<http://lbbe-shiny.univ-lyon1.fr/guts-shinyapp/>

Video-03

<https://sites.google.com/view/preditox2020>



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General information

Reference : UMR5558-SANCHA-002

Workplace : VILLEURBANNE

Date of publication : Friday, March 22, 2019

Type of Contract : FTC Technical / Administrative

Contract Period : 12 months

Expected date of employment : 3 June 2019

Proportion of work : Full time

Remuneration : Between 2423€ and 2612€ of monthly gross pay according to experience

Desired level of education : Higher than 5-year university degree

Experience required : 1 to 4 years

Missions

The main mission lies in revisiting species sensitivity distribution (SSD) methods in the context of environmental risk assessment of pesticides on non-target terrestrial plants (NTTP). In particular, the applicant will have to:

- Select appropriate non-linear models to obtain statistically robust estimates of concentrations for which there is a 50% effect on the species tested (ER50);
- Define an SSD procedure to include censored ER50 estimates (ie defined as intervals and / or unbounded values) allowing to statistically estimate hazardous concentrations for 5% of species (HC5), covering the full range of sensitivities displayed in studied sets of NTTP;
- Apply this SSD procedure to construct an applicable decision tree for risk assessment based on case studies;
- Design a user-friendly user interface, compatible with good laboratory practices (GLP) applicable to computer systems, making it easy to implement the decision tree and the SSD procedure.

Faites connaître cette offre !

URL Courte :

<http://bit.ly/2WkhnNH>

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 Partager

A dark blue, irregularly shaped graphic with a splatter effect, containing the text "Thank you!" in white. The graphic has a rough, hand-painted appearance with various shades of blue and white splatters around its edges. The text is centered within the dark blue area.

Thank you!