

Inclusion of Toxicity Information into Identification of Alerting Chemicals

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Our Concerns

How to include **toxicity** information into the identification of **alerting** chemicals?

- Prioritising: **Screening**
- Risk assessment: As **precise** as possible

How to **obtain** toxicity **information**?

- **Experimental** data
- **Predictive** models

Toxicity Prediction

- Many **different organisms** and **endpoints**
 - **Ecotoxicology** vs. **human** health
 - Toxicity depends on **mode of action**
-
- **Quantitative** models **limited, restricted**
 - **Alternatives:**
 - **Structural alerts, read-across**
 - **Advances approaches** to be explored

Baseline Toxicity

- Reversible **damage** of cell **membrane**
- Toxicity driven by **hydrophobicity**
- **Quantitative** estimation **possible**

Linear relationship hydrophobicity / toxicity (logarithmic scale)

$$\log EC_{50} = A \cdot \log K_{ow} + B$$

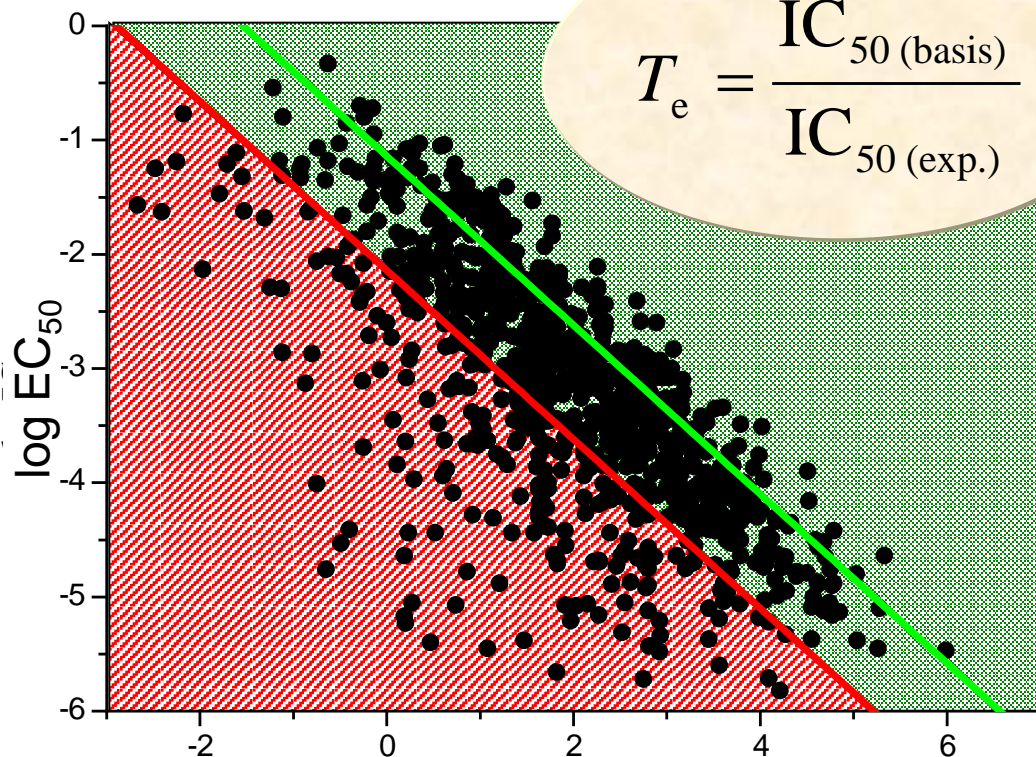
Excess Toxicity

- Significantly **higher** toxicity than estimated by **baseline** toxicity equation
- **Reactive** or **specific** modes of action

Note: High toxicity ~ low EC₅₀

Baseline and Excess Toxicity

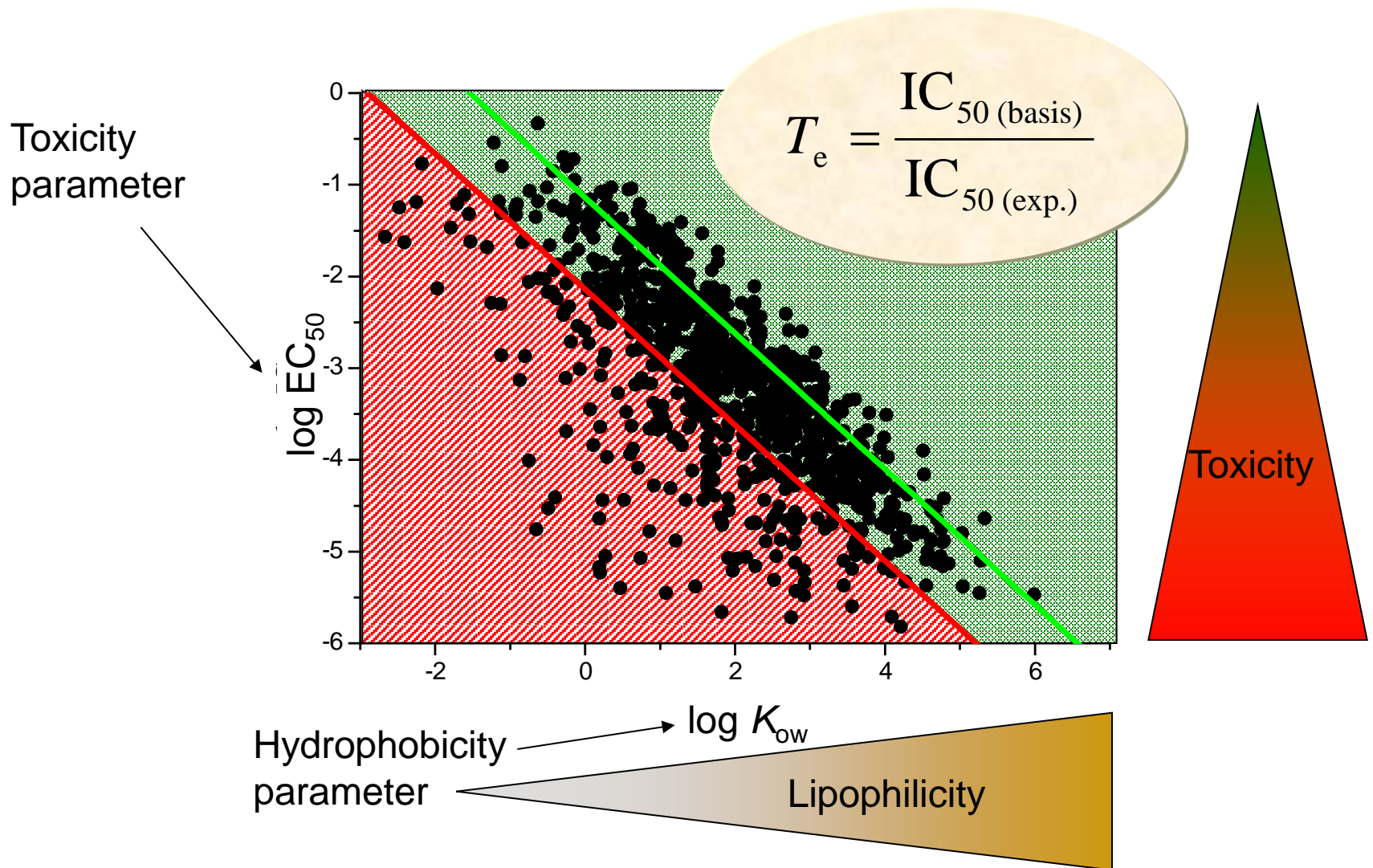
Toxicity
parameter



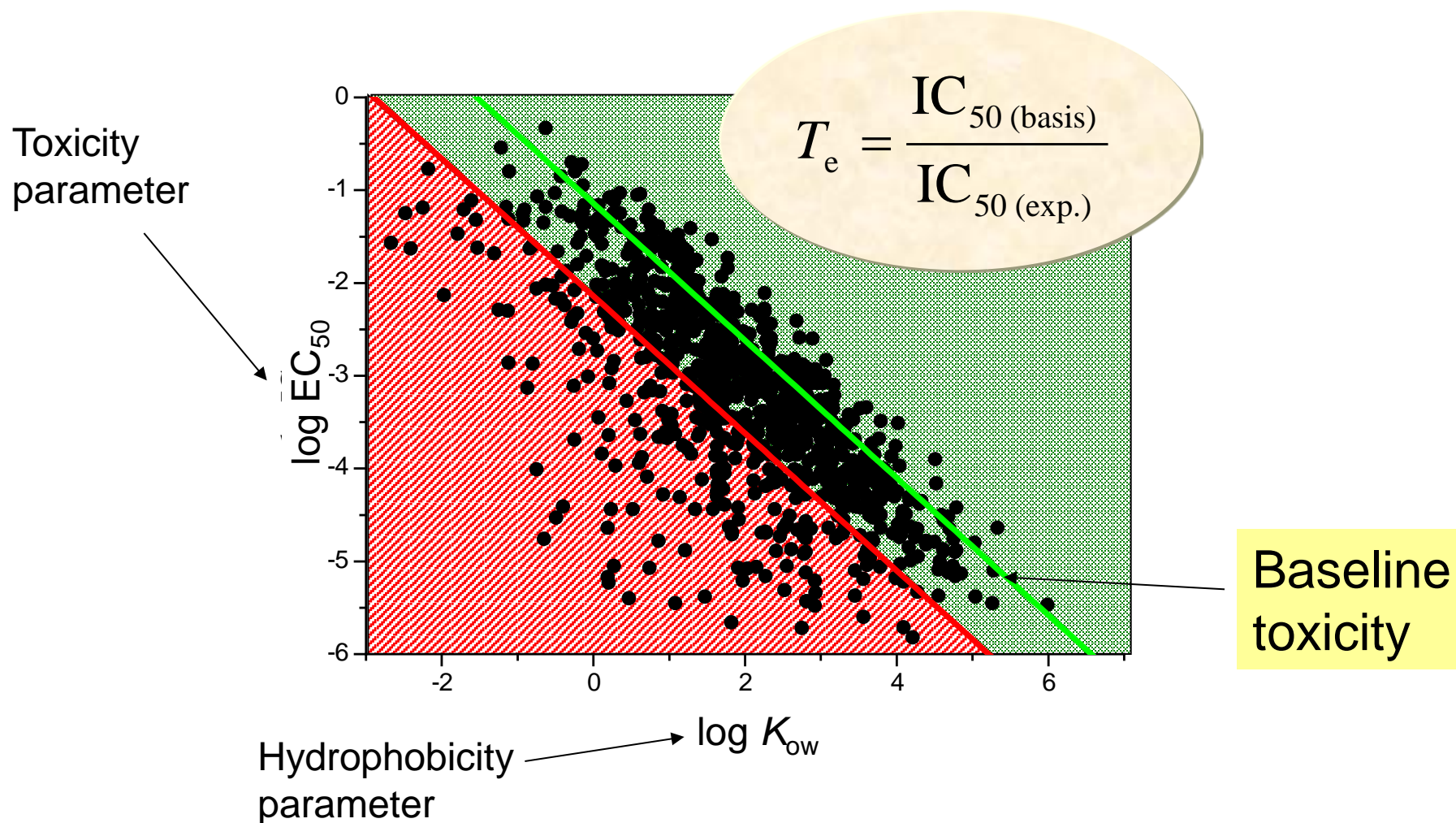
Hydrophobicity
parameter

$\log K_{ow}$

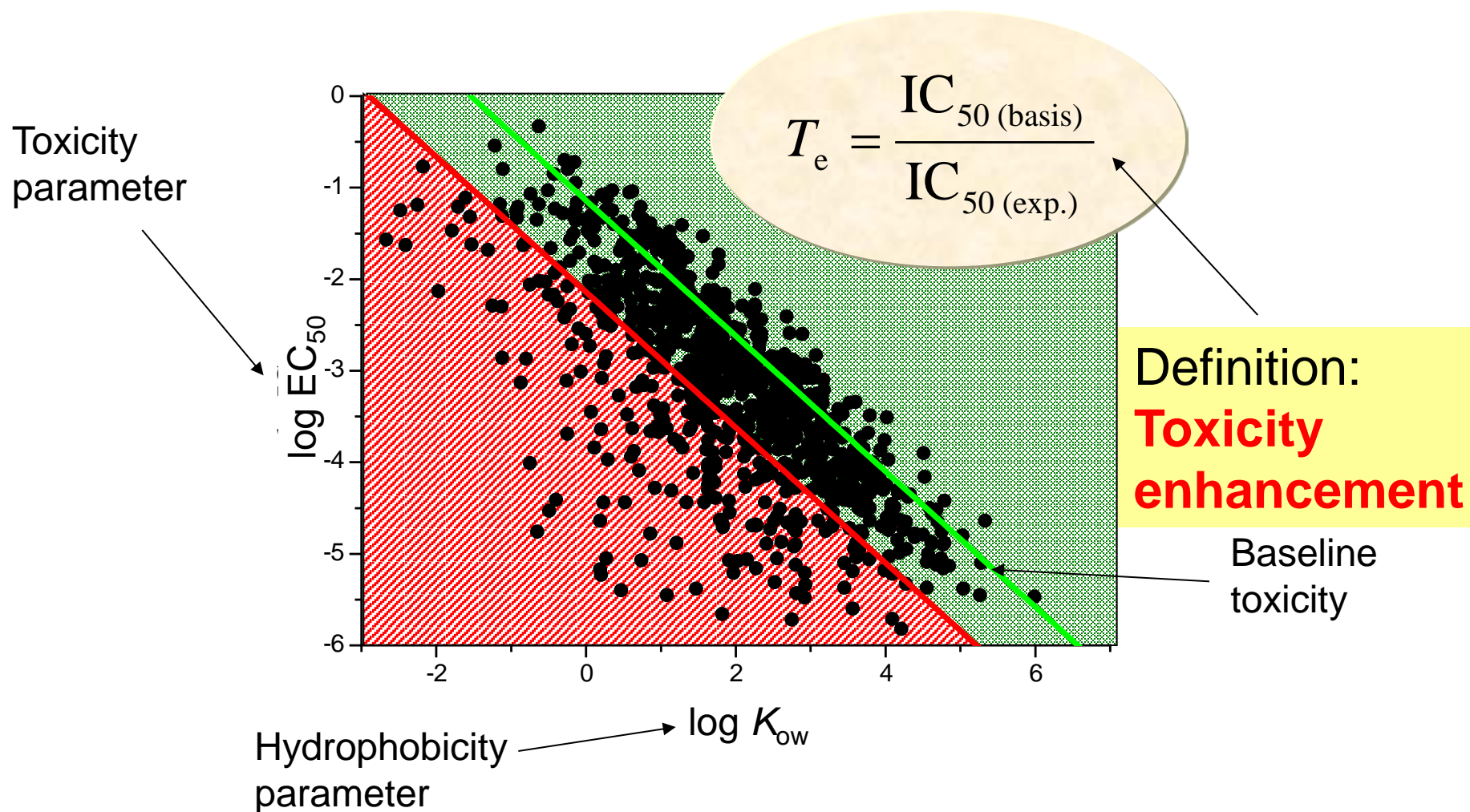
Baseline and Excess Toxicity



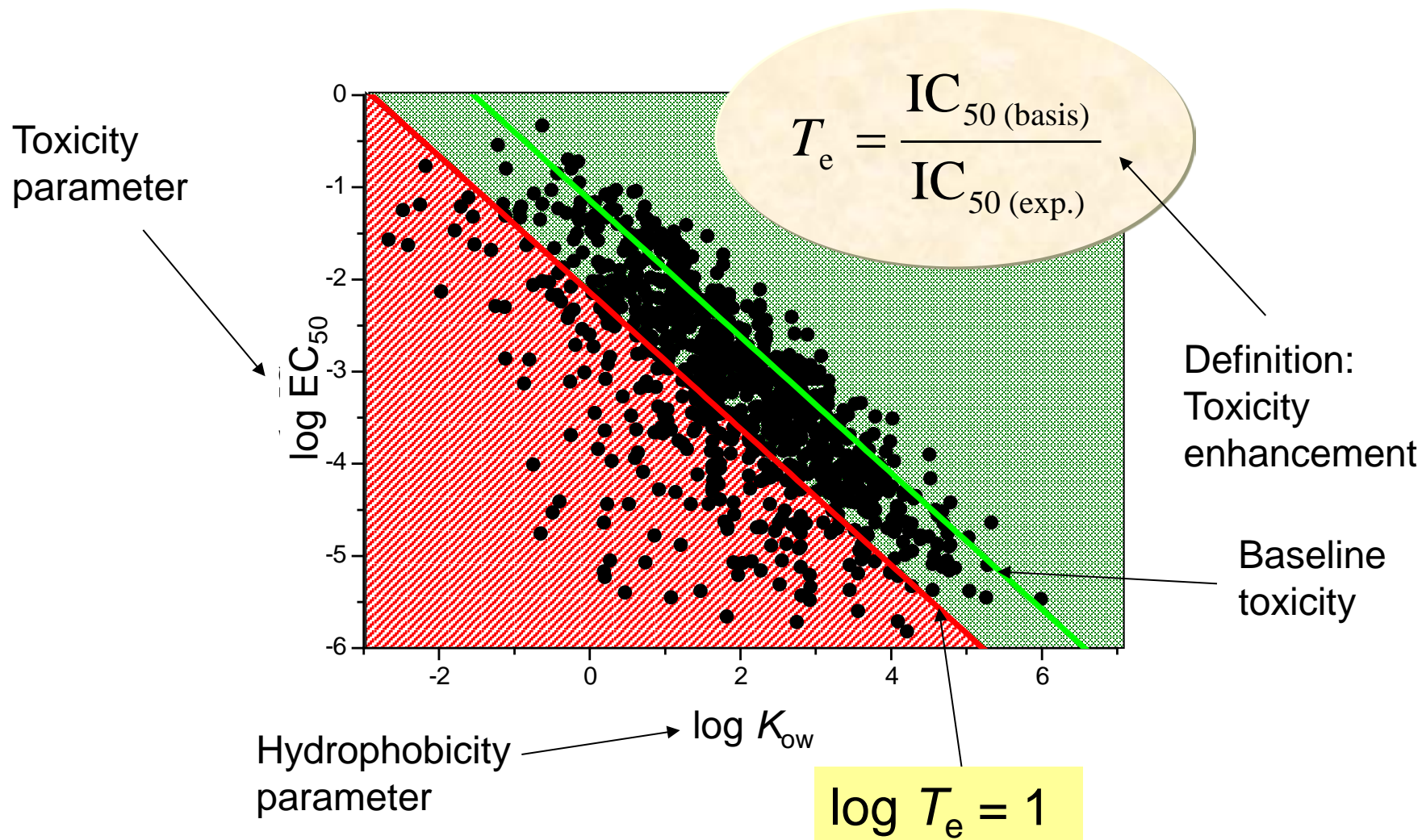
Baseline and Excess Toxicity



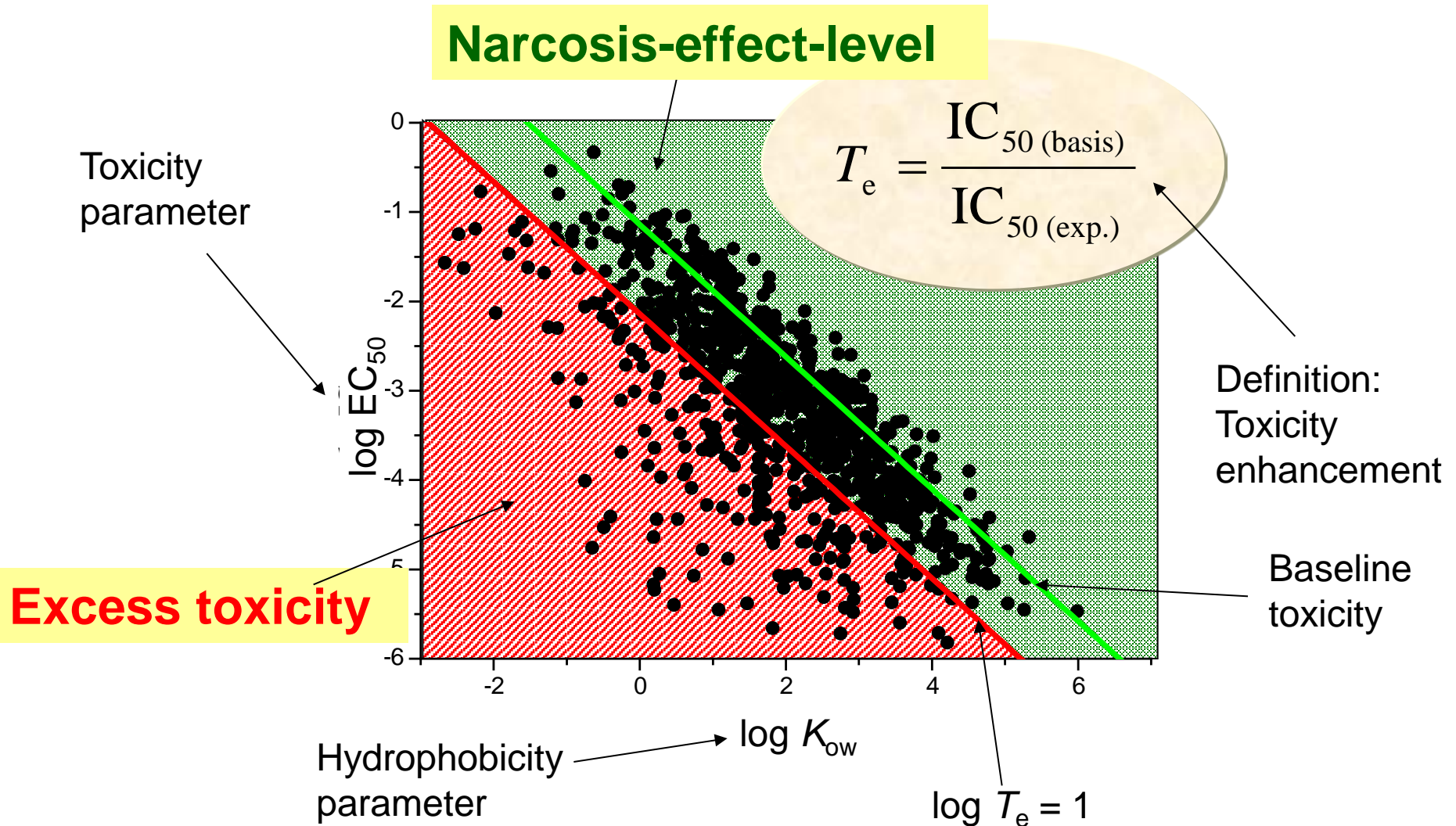
Baseline and Excess Toxicity



Baseline and Excess Toxicity



Baseline and Excess Toxicity



Structural Alerts for Predicting Aquatic Excess Toxicity

Hypothesis

Molecular **reactivity** related to **excess** toxicity

General Approach

Reactive **functional groups** as basis for excess toxicity model

New Structural Alert Model for Fish

23 Structural Alerts for Excess Toxicity

T_e (Toxicity Enhancement) Threshold

- $\log LC_{50} \leq \log LC_{50, \text{baseline}} - 1$

Training Set

- Validated LC_{50} (fathead minnow) for **693 cpds**

Data Sources

- **Duluth** database ($n = 526$)
ECOTOX/Aquire database ($n = 170$)

Structural Alert Model for Fish: Model Performance

Fathead minnow $n = 693$

Narcosis-Effect-Level : No structural alert
Good prediction power (95%)

Excess Toxicity: At least 1 of 23 structural alerts
Good prediction power (90%)

EINECS Compounds: Structural Alerts for Excess Toxicity

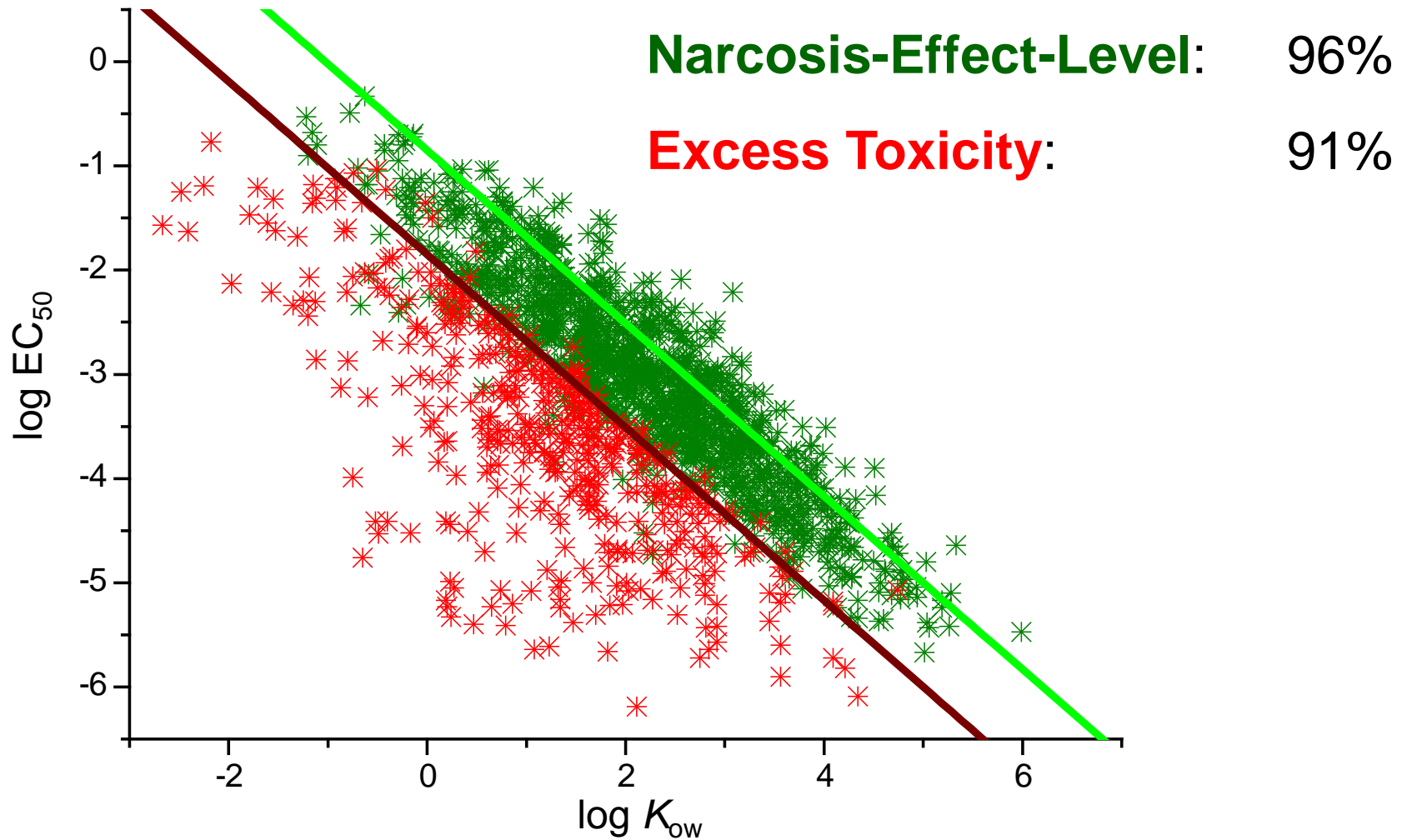
Fathead minnow:

Acute Toxicity, 96-h exposure, log LC₅₀ (*n* = 72520)

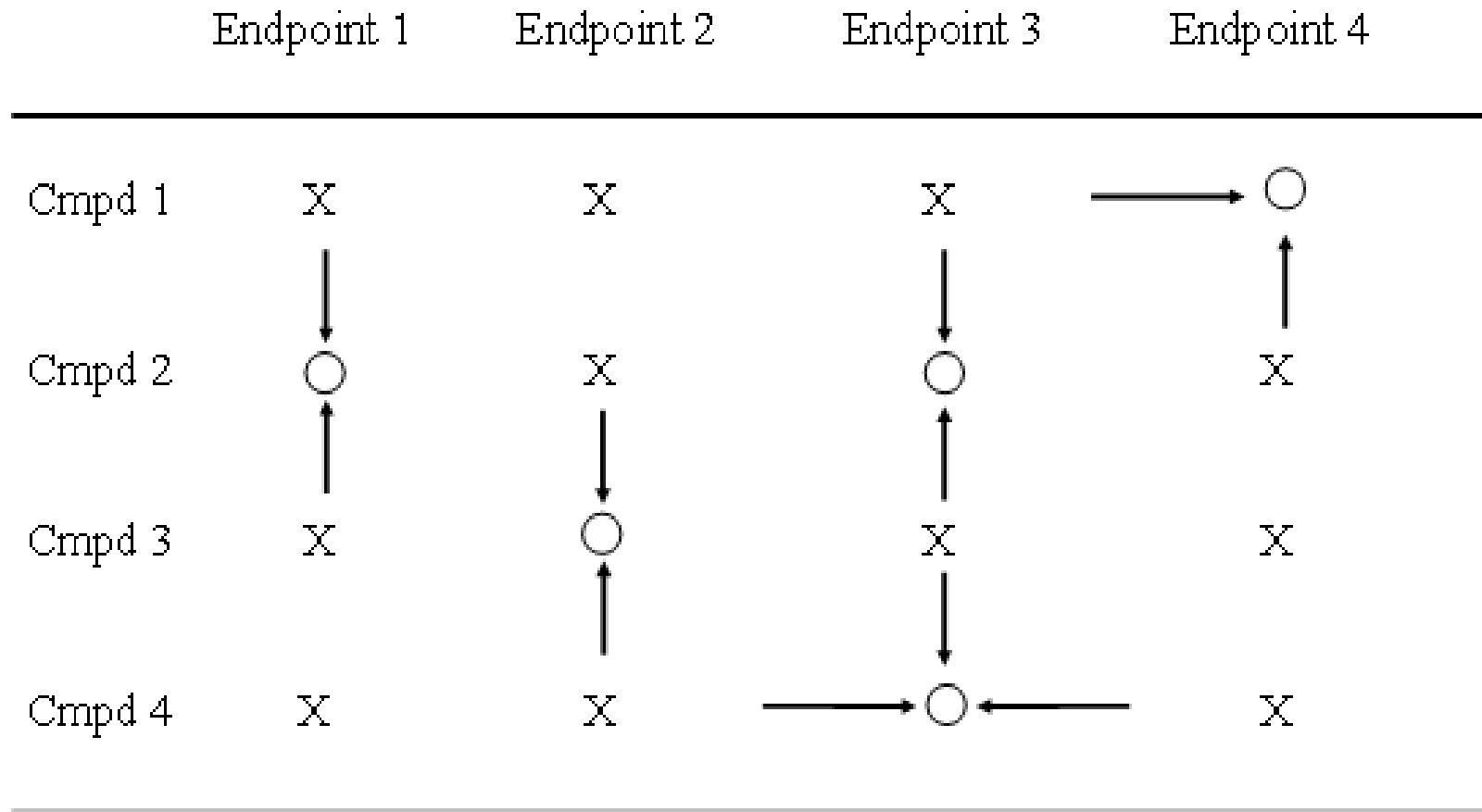
<i>less strict dom.</i>	<i>all cpds</i>	<i>not strictly out of domain</i>	<i>in domain</i>
Alert	12630	7954	3153
No alert	59890	35313	16065
No result	0	29253	53302

- Prediction for (only) 60%,
with **49% narcosis-level**, and **11% excess-toxic**

Predictivity SA model *T.pyriformis*



Read-across



Fish Toxicity: Read-across

Fathead Minnow:

Acute Toxicity, 96-h exposure, log LC₅₀ (*n* = 692)

	<i>n</i> valid	<i>r</i> ² LOO	<i>q</i> ² LOO	<i>rms</i> LOO	<i>bias</i>	<i>mne</i>	<i>mpe</i>
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Similarity Threshold

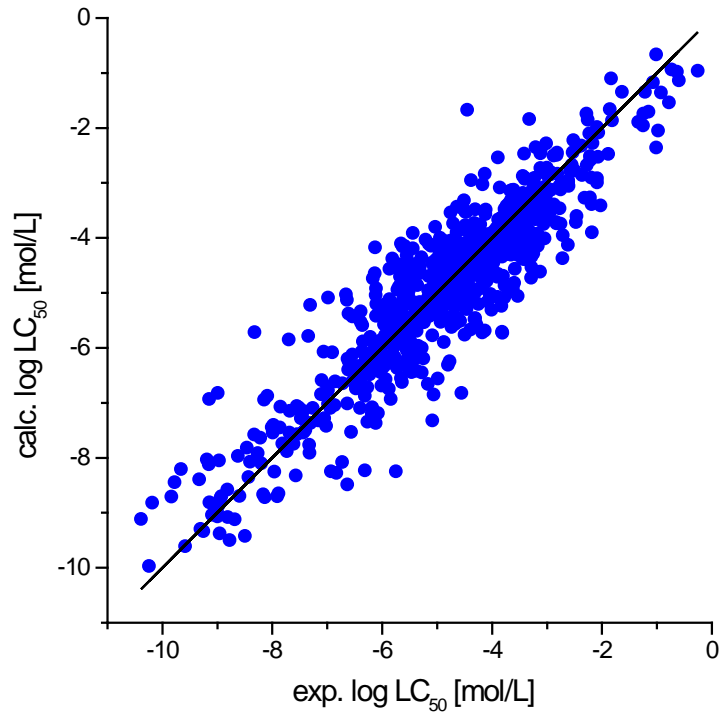
No	692	0.73	0.72	0.73	0.03	-2.31	4.29
Moderate	419	0.78	0.76	0.60	0.00	-2.92	3.99
High	230	0.84	0.87	0.43	-0.01	-1.34	1.73

Schüürmann et al., *EST* 2011

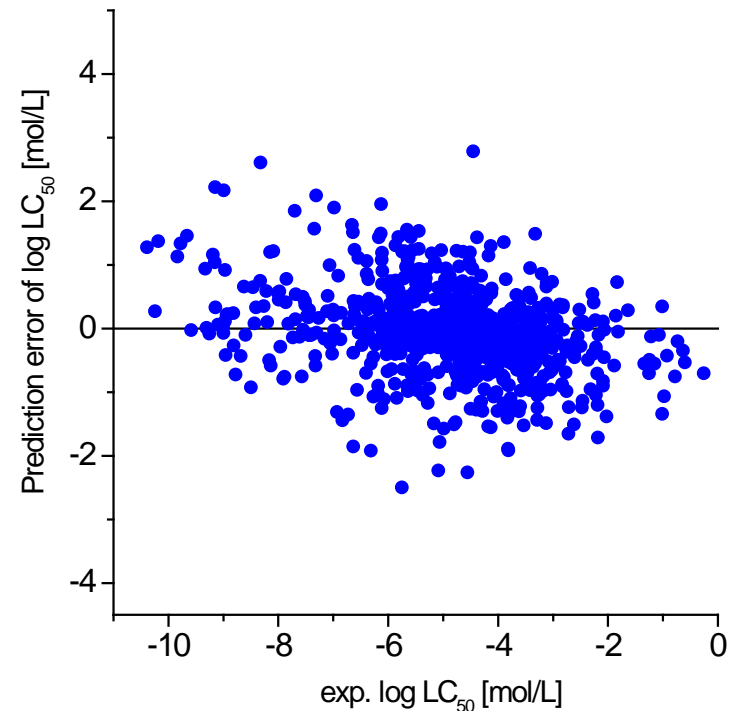
Daphnid Toxicity: Quantitative Model

Log LC₅₀ (mol/L)

Calculated vs experimental



Residues vs experimental



$n = 759$ $r^2 = 0.83$ $q^2 = 0.83$ $rms = 0.68$ $bias = -0.04$

Kühne et al., *Mol. Inf.* 2013

Daphnid Toxicity: Qualitative Model

- **Excess toxicity prediction**
 $n = 50$, 47 of them (94%) correctly classified

- **Screening level classification** ($n = 271$)

Narcosis level

Sensitivity 86%

Predictivity 82%

Excess toxicity

Sensitivity 78%

Predictivity 83%

Concordance 82%

- **Out of domain** $n = 285$ (21%)

Fish Toxicity: Consensus Modeling for Excess Toxicity Probability

Read-across; ECOSAR; Structural alerts

<i>Model T_e pred.</i>	<i>N</i>	<i>Pred. prob. excess</i>	<i>exp. excess</i>
+ + +	75	100%	95%
+ + -	29	80%	21%
+ - +	21	98%	76%
- + +	38	99%	89%
+ - -	33	22%	12%
- + -	48	27%	8%
- - +	23	83%	83%
- - -	427	3%	3%

QSAR Software

- ChemProp (UFZ)

The screenshot displays the ChemProp software interface. On the left, a list of 15 compounds is shown, including 2-PROPYLPYRIDINE, 2-CHLOROTOLUENE, 4-AMINO-2-NITROPHENOL, and others. Below this is a 3D ball-and-stick model of a molecule. The main window is titled 'ChemProp Methods selection' and shows a 'RESULTS: Read across 3' table. The table has columns for 'STRUCTURES', 'RESULTS: Read across 3', and 'Molar weight range'. The results table contains the following data:

STRUCTURES	RESULTS: Read across 3	Molar weight range	
1: 2-PROPYLPYRIDINE	1.5	3	-1 n.a., 1 >1250
2: 2-CHLOROTOLUENE	1.6	3	3 -4.8, 1 >1000
3: 4-AMINO-2-NITROPHENOL	2.1	3	-2 <-4.8, 2 >4.9
4: PROPYL ACETATE	2.2	3	-1 <-4.5, 1 >11
5: 2,4,6-TRINITROPHENOL	2.2	3	3: has 0, 0, 0
6: 1,4-DINITROBENZENE	2.2	3	2: has 0, 1 >0.8, 0 >1
7: 2,4-DINITROPHENOL	2.3	3	3: -1.15 ... 2.42 (AA), -2 <-1.75, 2 >7.62
8: 4-AMINOPHENOL	2.3	3	-1 <-2.9, 1 >8.30

Below the table, there is a section for 'Read-across (Schüürmann et al)' with a description: 'is mol L is calculated from the toxicity enhancement T_e and from the toxicity LC_{50} (narc) by $Log LC_{50} = log LC_{50} (narc) - log T_e$. $Log LC_{50}$ (narc) is estimated from $log K_{ow}$ by $Log LC_{50} (narc) = -0.85 \cdot log K_{ow} - 1.41$ '.

- ToxTree, OECD Toolbox, EPISUITE
- Commercial software (e.g. DEREK, Multicase, ...)

Beyond LC₅₀

Nominal concentrations do **not** reflect amount of chemical at **target site**

- Physiologically based pharmacokinetics (**PBPK**) or PBTK
- Replace concentrations by **activities**

Response from **cell** up to **ecosystem**

- Adverse Outcome Pathways (**AOP**)

Integrating Approaches

- Integrated Testing Strategies (**ITS**)



- Integrating knowledge across **human** and **environmental** risk assessment



Exposome

Take Home Messages

- **Toxicity data limited**
- **Quantitative models typically restricted to Mode of Action**
- **Structural alerts useful screening tool**
- **QSAR models complemented by read-across**
- **Consensus of different approaches preferable**
- **Crucial: Applicability domain**

UFZ Team

Head: Prof. Dr. Gerrit Schüürmann
Dr. Norbert Ost, Ralf-Uwe Ebert,
many more ...

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