



Weight of Evidence by Conditional Probabilities: A Bayesian network model for predicting fish acute toxicity based on fish embryo testing

[Jannicke Moe](#), Anders L. Madsen, Raoul Wolf, Kristin A. Connors, Jane M. Rawlings,
Scott E. Belanger, Wayne G. Landis, T. Braunbeck, M. Embry, K. Schirmer, S. Scholz, Adam D. Lillicrap

**Session 4.01: Artificial Intelligence Approaches in Environmental Risk Assessment:
Bayesian Networks, Machine Learning and Predictive Modelling**

Presentation ID: 4.01.08

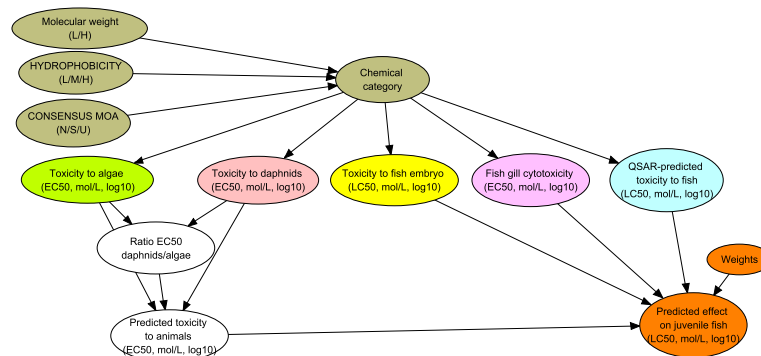


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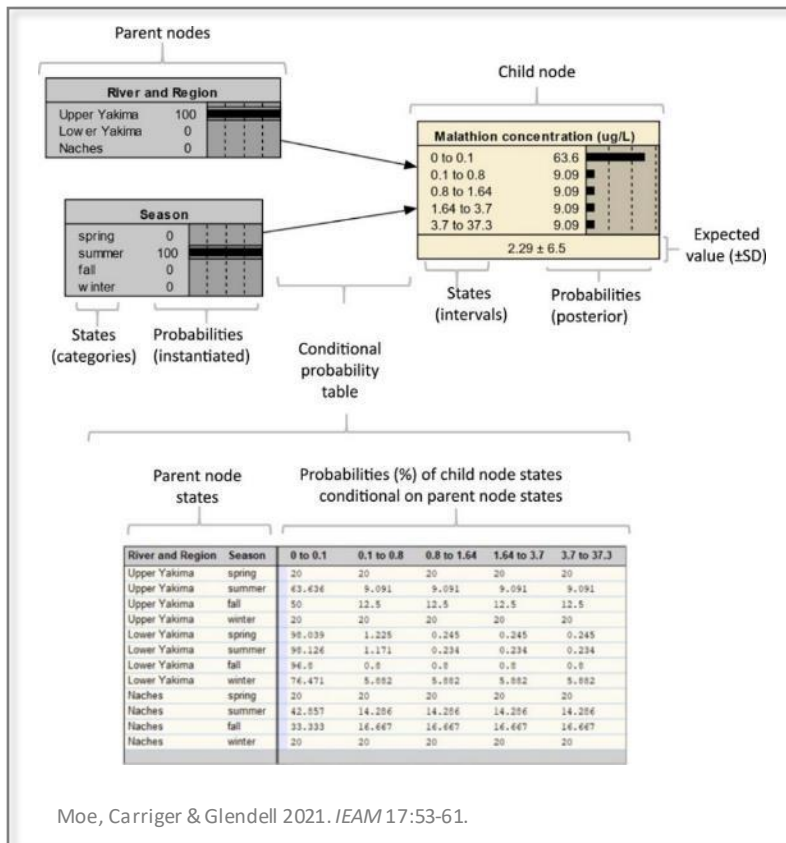


Highlights: We have developed a Bayesian network model which ...

- ✓ integrates fish **embryo toxicity data** with other information on physical, chemical and toxicological properties of substances
- ✓ represents a **quantitative weight-of-evidence** approach
- ✓ predicts the **correct fish acute toxicity** intervals for most of test substances
- ✓ is **publicly available** for demonstration and testing in a web interface



What is a Bayesian network?



- A joint probability distribution among variables (nodes) in a graphical format
- Relationships (arrows) are quantified by Conditional Probability Tables (CPTs)
- The nodes usually have discrete states
- The nodes and arrows form a directed acyclic graph; no feedback loops
- Many recent applications in ERA

Special Series: Applications of Bayesian Networks for Environmental Risk Assessment and Management



Increased Use of Bayesian Network Models Has Improved Environmental Risk Assessments

S Jannicke Moe, John F Carriger, Miriam Glendell

First published: 18 November 2020 | <https://doi.org/10.1002/ieam.4369>

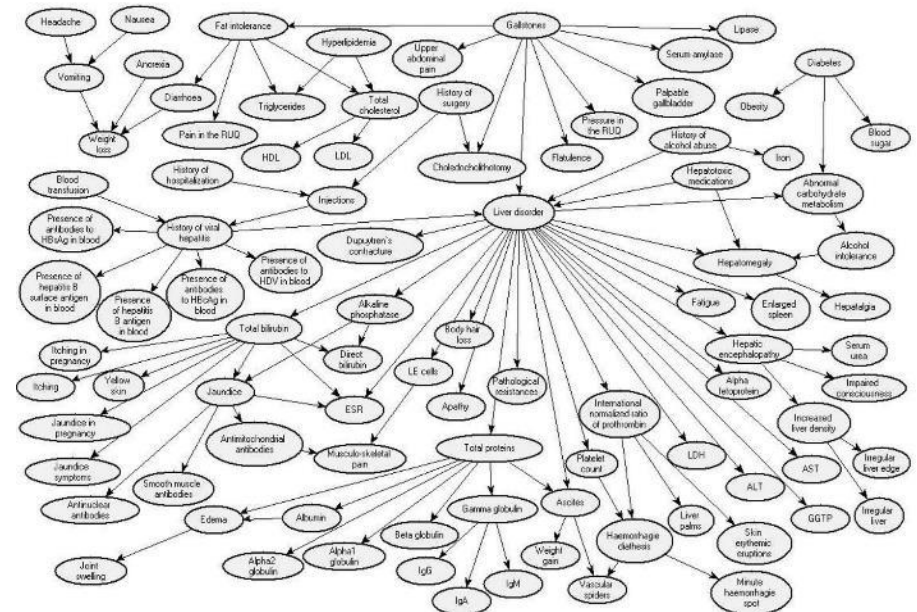
Moe, Carriger & Glendell 2021. *IEAM* 17:53-61.

BNs can use Artificial Intelligence for learning model structure and parameter values



- Machine learning (ML) methods require large data sets
- Toxicity and exposure data are often limited
- “Unlike most ML methods, BNs allow for direct **expert knowledge** input, to control the direction and existence of edges between nodes”
- “BNs fill an important gap in the ML world, **bridging the divide** between simple models without probability information, and computationally heavy and data-hungry methods”

After <https://medium.com/eliiza-ai/bayesian-networks-combining-machine-learning-and-expert-knowledge-into-explainable-ai-efaf6f8e69b>



<https://www.norwegiancreations.com/2018/09/artificial-intelligence-bayes-network/>

A BN model for predicting acute **fish toxicity** from **fish embryo toxicity** & other information

Environmental Modelling and Software 126 (2020) 104655

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journal homepage: <http://www.elsevier.com/locate/envsoft>

Development of a hybrid Bayesian network model for predicting acute fish toxicity using multiple lines of evidence

S. Jannicke Moe^{a,*}, Anders L. Madsen^{b,c}, Kristin A. Connors^d, Jane M. Rawlings^d, Scott E. Belanger^d, Wayne G. Landis^e, Raoul Wolf^e, Adam D. Lillicrap^a

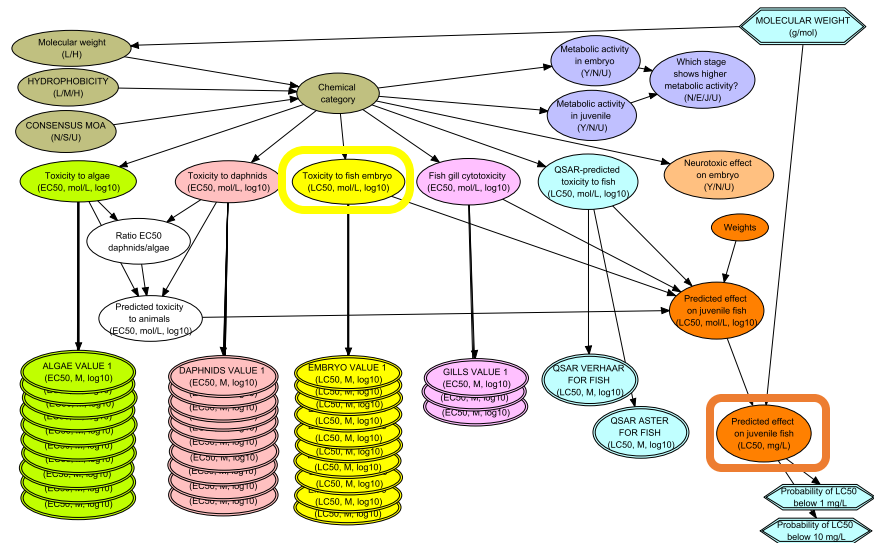
Integrated Environmental Assessment and Management — Volume 16, Number 4—pp. 452–460

452 Received: 5 September 2019 | Returned for Revision: 21 November 2019 | Accepted: 20 February 2020

Decision Analysis

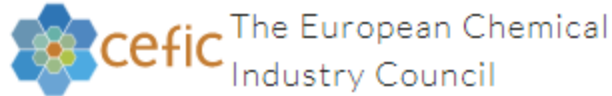
Evaluation of a Bayesian Network for Strengthening the Weight of Evidence to Predict Acute Fish Toxicity from Fish Embryo Toxicity Data

Adam Lillicrap,^{a,*} S Jannicke Moe,[†] Raoul Wolf,[†] Kristin A Connors,[‡] Jane M Rawlings,[‡] Wayne G Landis,[§] Anders Madsen,^{||} and Scott E Belanger,[‡]



- Aim: replace the use of juvenile fish (OECD 203) with fish embryos (OECD 236) in toxicity testing
- Request: «strengthen the weight of evidence for fish embryo data»
- (More background information: presentation 1.01.07)

The SWiFT project (2020-2023)



Strengthening **Weight** of evidence for FET data to replace acute **Fish Toxicity** (SWiFT)

Researcher Team:

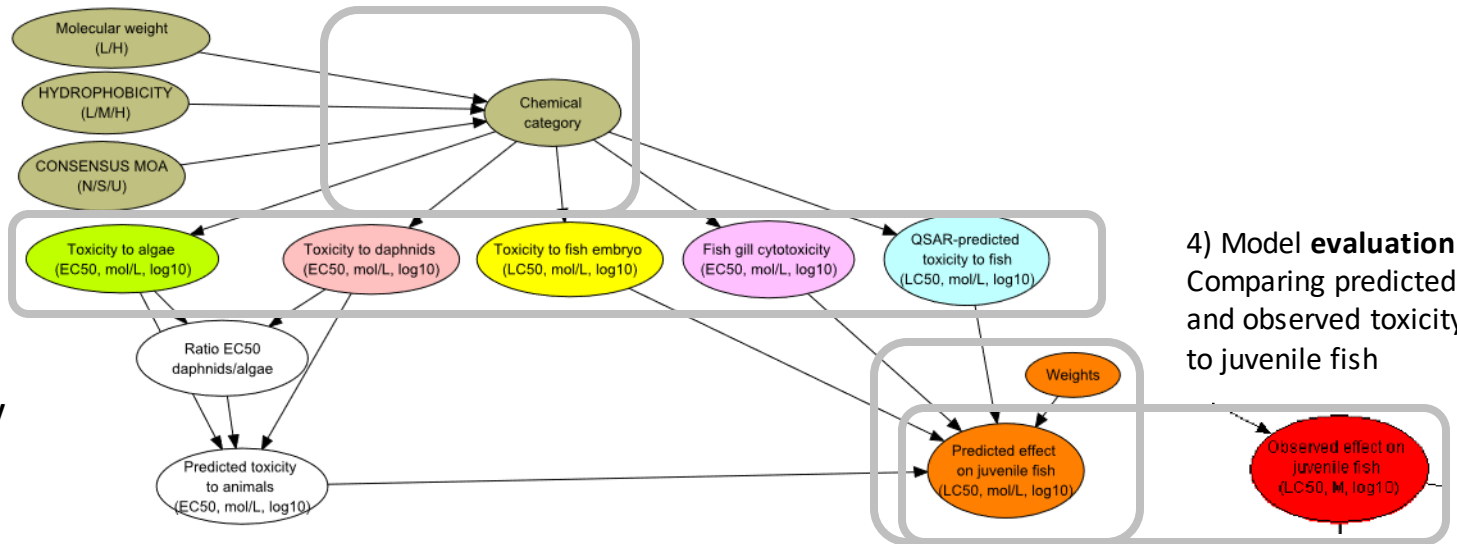
Adam Lillicrap (PI, WP2), Kristin Connors (WP1), Jannicke Moe (WP3), Anders Madsen (WP4), Raoul Wolf, Thomas Braunbeck, Kristin Schirmer, Michelle Embry, Scott Belanger, Stefan Scholz

Advisory Team:

Noemie Croze, Christopher Faßbender, Sylvia Gimeno, Marlies Halder, Sarah Hughes, Joop de Knecht, Mark Lampi, Wayne Landis, Teresa Norberg-King, Martin Paparella, Audrey Pearson, Eleonora Simonini, Marta Sobanska, Susanne Walter-Rohde

Main properties of the BN model, run for one chemical substance

- 0) Enter info on physical & chemical properties of the substance
→ define chemical category
→ get **prior probability** of toxicity for each Line of Evidence (LoE)



1) Enter toxicity values (= **evidence**) for one or more Lines of Evidence

2) combine with priors
→ **posterior probability** of toxicity for each LoE

4) Model **evaluation**: Comparing predicted and observed toxicity to juvenile fish

3) **Integration** and prediction: Weighting of each LoE

How can our BN be used in a WoE approach?

Guidance on the use of the weight of evidence approach in scientific assessments

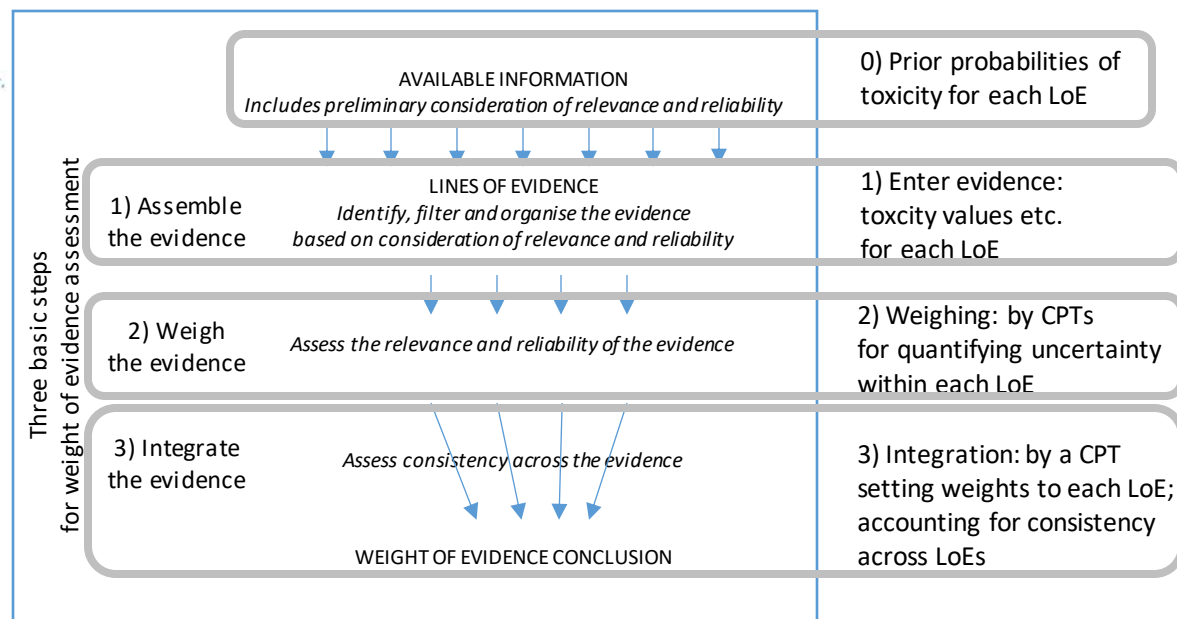
EFSA Scientific Committee, Anthony Hardy, Diane Benford, Thorhallur Halldorsson, Michael John Jeger, Helle Katrine Knutsen, Simon More, Hanspeter Naegeli, Hubert Noteborn ... See all authors

First published: 03 August 2017 | <https://doi.org/10.2903/j.efsa.2017.4971> | Citations: 47

Our BN-WoE should be

- **consistent** with WoE approaches recommended for regulatory frameworks (ECHA, EFSA, US EPA)
- **quantitative**
- **intuitive**
- **flexible**

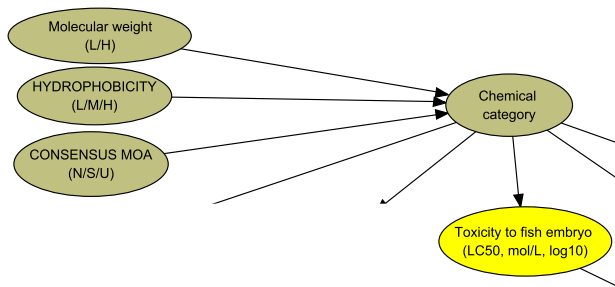
Main steps of the BN model:



0) Prior probability of toxicity: existing toxicity data grouped by chemical category

Definition of chemical categories:

- Molecular weight (**L**ow/**H**igh)
- Hydrophobicity (**L**ow/**M**edium/**H**igh)
- Mode of action (**N**arcotic/**S**pecific/**U**nspecified)



Category «LLU»:

Wide probability distribution
= high uncertainty

Category «LMS»:

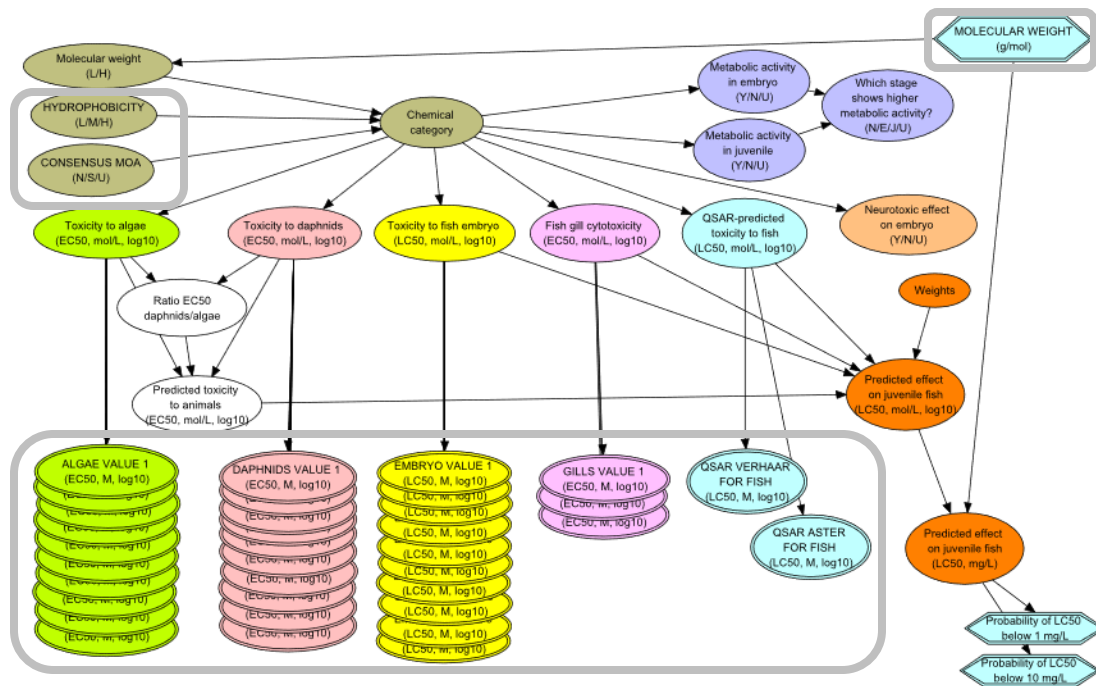
Narrow probability distribution
= low uncertainty

- Training data set: 237 substances (Moe et al. 2020)
- Current method: count of observations
- In development: hierarchical Bayesian estimation (Presentation 1.01.07)



Chemical c...	LLN	LLS	LLU	LMN	LMS	LMU	LHN	LHS	LHU	HLN	HLS	HLU	HMN	HMS	HMU	HHN	HHS	HHU	
-inf -9	9.998001E-9	9.998005E-9	9.99801E-5	9.997994E-9	9.996002E-9	9.997001E-9	9.994993E-9	9.994004E-9	9.994004E-9	9.998005E-9	9.998005E-9	9.994004E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9
-9 -6	0.017003	0.086403	0.166633	0.217348	0.121164	0.235629	0.333167	0.9994	9.994004E-9	0.09884	0.09884	9.994004E-9	0.09884	0.09884	0.09884	0.09884	0.09884	0.09884	0.09884
-6 -5	0.064613	0.185148	0.208292	0.434696	0.605818	0.420443	0.666334	9.994004E-9	0.9994	0.171068	0.171068	9.994004E-9	0.171068	0.171068	0.171068	0.171068	0.171068	0.171068	0.171068
-5 -4	0.265253	0.469042	0.231608	0.173878	0.272618	0.142814	9.994993E-9	9.994004E-9	9.994004E-9	0.264206	0.264206	0.9994	0.264206	0.264206	0.264206	0.264206	0.264206	0.264206	0.264206
-4 -3	0.319664	0.135775	0.166633	0.130409	9.996002E-9	0.142814	9.994993E-9	9.994004E-9	9.994004E-9	0.220488	0.220488	9.994004E-9	0.220488	0.220488	0.220488	0.220488	0.220488	0.220488	0.220488
-3 -0	0.333267	0.123432	0.166633	0.04347	9.996002E-9	9.997001E-9	9.994993E-9	9.994004E-9	9.994004E-9	0.245198	0.245198	9.994004E-9	0.245198	0.245198	0.245198	0.245198	0.245198	0.245198	0.245198
0 -inf	9.998001E-9	9.998005E-9	9.99801E-5	9.997994E-9	9.996002E-9	9.997001E-9	9.994993E-9	9.994004E-9	9.994004E-9	9.998005E-9	9.998005E-9	9.994004E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9	9.998005E-9
Experience	294	81	24	23	33	7	3	2	2	0	0	1	0	0	0	0	0	0	0

1) Assemble the evidence



Enter evidence for one substance:

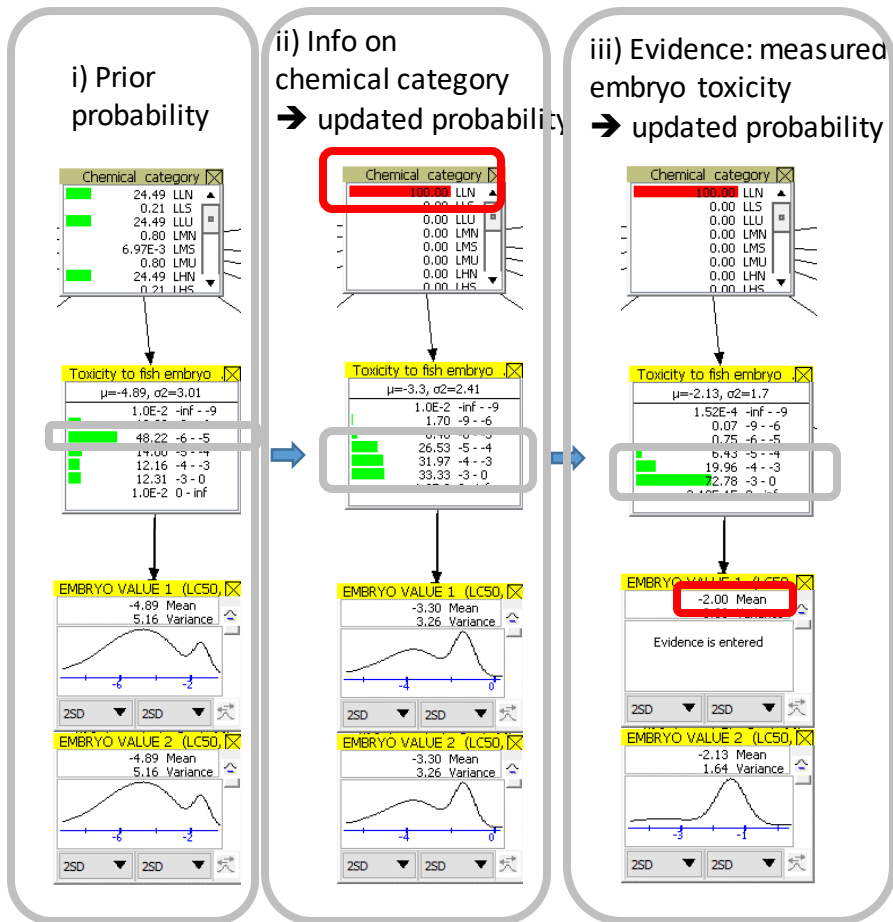
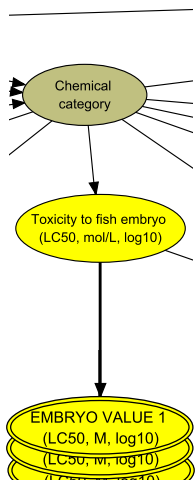
- Molecular weight (g/mol)
- Hydrophobicity (L/M/H)
- Mode of action (N/S/U)
- Algae (EC50, mg/L)
- Daphnids (EC50, mg/L)
- Embryo (LC50, mg/L)
- Gill cytotoxicity (EC50, mg/L)
- (Neurotoxicity EC50)

Built-in data processing:

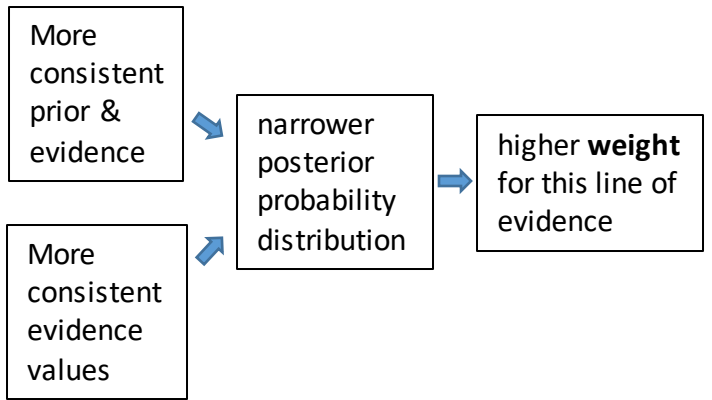
- conversion from mg/L to mol/L
- log10 transformation

2) Weigh the evidence

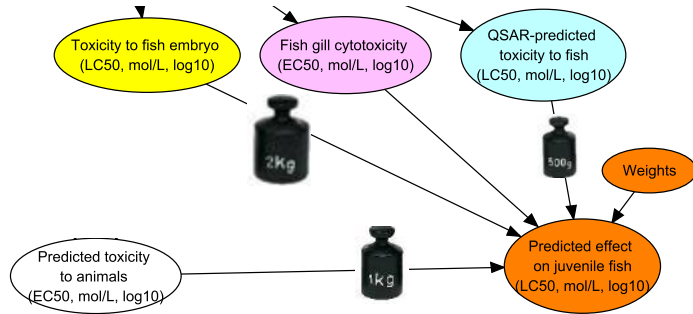
Example:
Embryo toxicity



iv) More evidence → updated probability etc.



3) Integrate & weight the lines of evidence



Predicted effect on juve. ✕		
$\mu=399.79, \sigma^2=181988.09$		
0.52	0 - 0.01	
0.32	0.01 - 0.1	
2.50	0.1 - 1	
16.27	1 - 10	
32.14	10 - 100	
22.55	100 - 1000	
25.72	1000 - inf	

«Correct prediction»

Observed effect on juve. ✕		
$\mu=355, \sigma^2=175621.97$		
1.13	0 - 0.01	
0.69	0.01 - 0.1	
1.97	0.1 - 1	
20.25	1 - 10	
33.68	10 - 100	
19.42	100 - 1000	
22.85	1000 - inf	

How to optimise the weighting of the LoEs?

- Optimisation of weights by **machine learning**?
 - Problem: requires $>10^6$ test cases
- We have max. 237 test substances
 - depending on the criteria for data selection
- Solution: define a limited set of scenarios with alternative weights
 - Evaluate model performance for alternatives
- Match the predicted vs. observed interval of LC_{50} for juvenile fish
 - Compare the most probable interval
 - How high precision is needed?

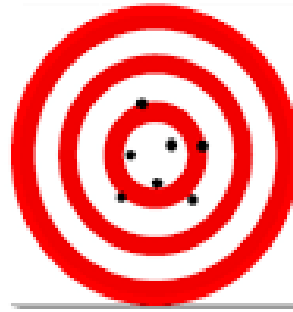
Precision vs. accuracy of the BN model predictions

HIGH ACCURACY,
HIGH PRECISION



Ideal situation

HIGH ACCURACY,
LOW PRECISION



Realistic situation:
aim of our BN model

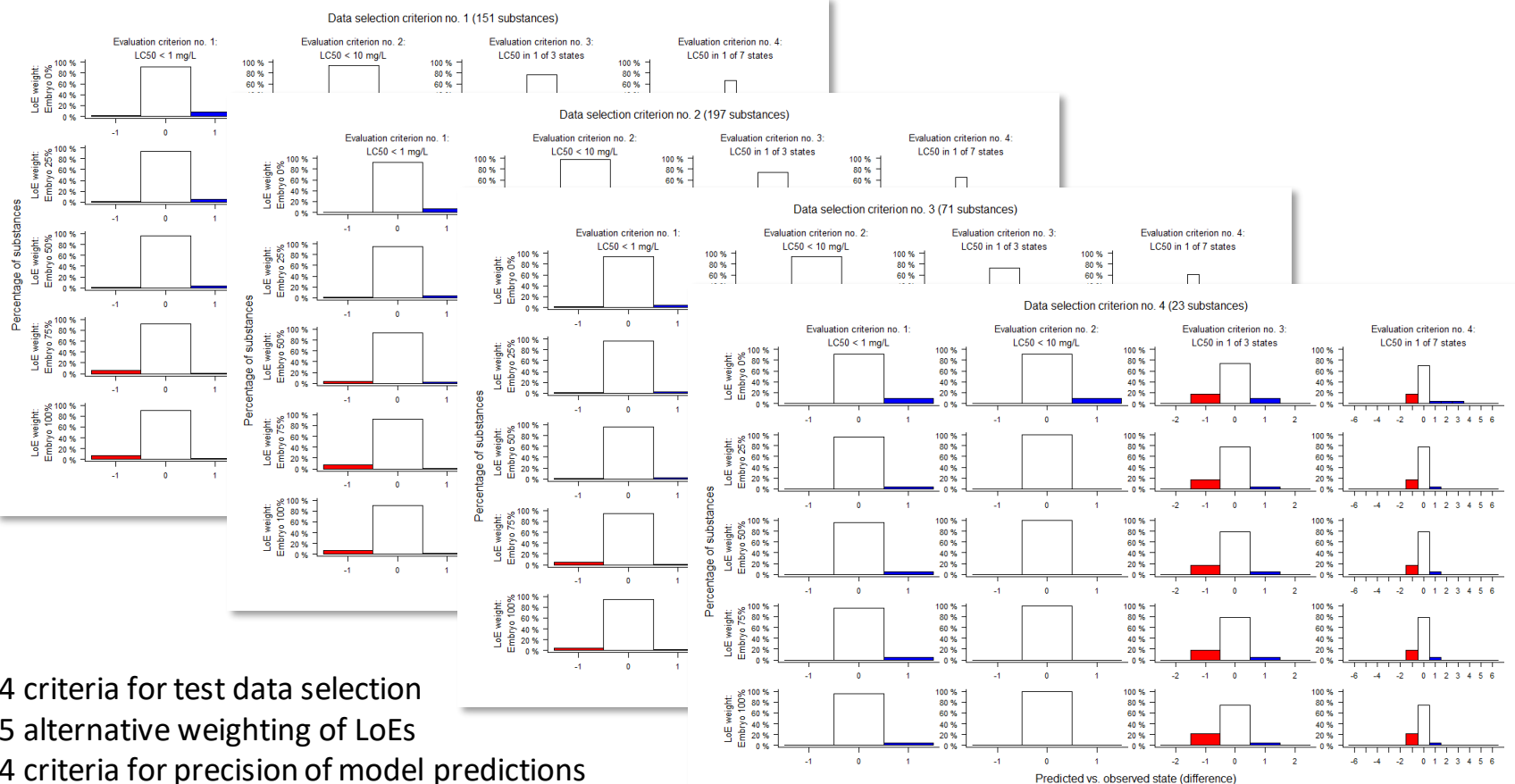
LOW ACCURACY,
HIGH PRECISION



Common situation:
point estimate
ignoring uncertainty

«It's better to be **roughly right**
than **precisely wrong**»

Scenarios for model evaluation



- 4 criteria for test data selection
- 5 alternative weighting of LoEs
- 4 criteria for precision of model predictions

The model's success rate depends on the required **precision** of predicted toxicity

Example:
1 substance

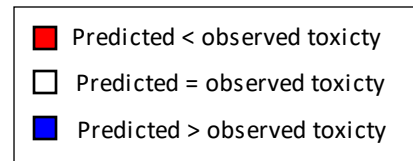
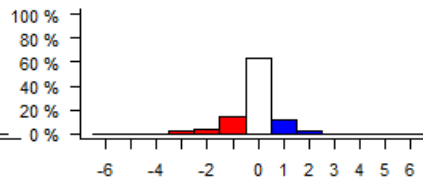
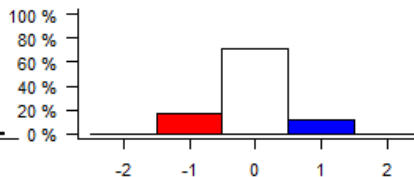
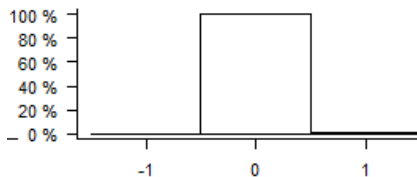
Predicted effect on juve. ✗	
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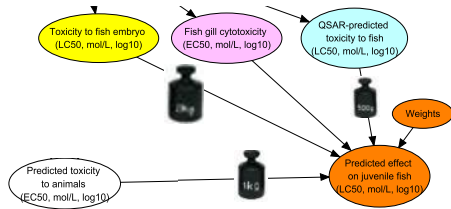
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Summary:
197 substances

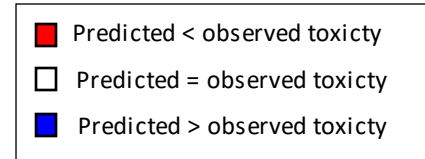
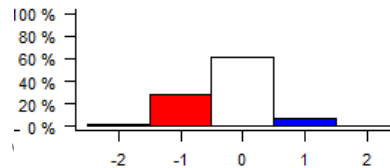
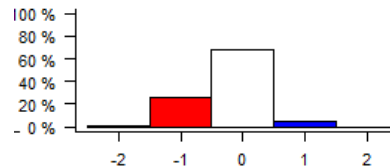
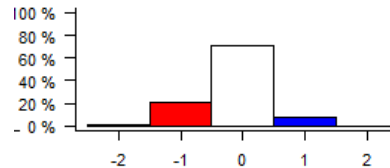
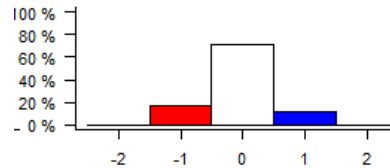
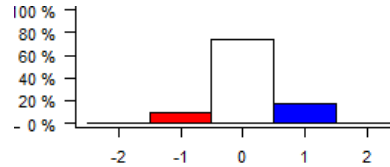


- Higher required resolution of predicted toxicity → lower success rate
- To be decided by stakeholders

The model's success rate depends on the weighting of lines of evidence



Weight scenario	Fish embryo	Algae & daphnids	QSAR fish
1	0 %	50.0 %	50.0 %
2	25 %	37.5 %	37.5 %
3	50 %	25.0 %	25.0 %
4	75 %	12.5 %	12.5 %
5	100 %	0 %	0 %



- Intermediate weight to Embryo (25-50%)
➔ highest accuracy
- Higher weight to Embryo
➔ more **underestimation** of acute fish toxicity

Conclusions

- Limited amount of test data (observed acute toxicity to juvenile fish)
 - ➔ limited possibility of machine learning for this BN
 - ➔ explore alternative approaches for training and testing
- The BN currently has a success rate of ca. 70-80 % of test substances, for predictions aggregated to 3 intervals (0-1, 1-10, 10-inf mg/L)
 - Balanced weighting of lines of evidence ➔ higher success
 - Lower resolution accepted for predicted toxicity ➔ higher success
- The BN-WoE approach is promising for supporting the use of fish embryo toxicity data instead of fish acute toxicity data.

Next steps

- Further exploration of model evaluation
 - Forthcoming larger test data set → explore machine learning methods?
 - Compare the whole distributions of predicted vs. observed toxicity
- Incorporate additional lines of evidence
 - Gill cytotoxicity; neurotoxicity; metabolic activity (presentation 1.01.07)
- Better account for uncertainty and variability in data
 - Hierarchical Bayesian regression (presentation 1.01.07)
- Further development of demonstration model with web interface
 - See Q&A field for updated information
 - Feedback appreciated!

Thanks for your attention!

Contact:

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More information on the SWiFT project:

<https://www.niva.no/swift>

<http://swift.hugin.com/>

or visit the NIVA online stand at SETAC Europe 2021

