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Decision Analysis

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

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ABSTRACT

The use of fish embryo toxicity (FET) data for hazard assessments of chemicals, in place of acute fish toxicity (AFT) data, has long been the goal for many environmental scientists. The FET test was first proposed as a replacement to the standardized AFT test nearly 15 y ago, but as of now, it has still not been accepted as a standalone replacement by regulatory authorities such as the European Chemicals Agency (ECHA). However, the ECHA has indicated that FET data can be used in a weight of evidence (WoE) approach, if enough information is available to support the conclusions related to the hazard assessment. To determine how such a WoE approach could be applied in practice has been challenging. To provide a conclusive WoE for FET data, we have developed a Bayesian network (BN) to incorporate multiple lines of evidence to predict AFT. There are 4 different lines of evidence in this BN model: 1) physicochemical properties, 2) AFT data from chemicals in a similar class or category, 3) ecotoxicity data from other trophic levels of organisms (e.g., daphnids and algae), and 4) measured FET data. The BN model was constructed from data obtained from a curated database and conditional probabilities assigned for the outcomes of each line of evidence. To evaluate the model, 20 data-rich chemicals, containing a minimum of 3 AFT and FET test data points, were selected to ensure a suitable comparison could be performed. The results of the AFT predictions indicated that the BN model could accurately predict the toxicity interval for 80% of the chemicals evaluated. For the remaining chemicals (20%), either daphnids or algae were the most sensitive test species, and for those chemicals, the daphnid or algal hazard data would have driven the environmental classification. *Integr Environ Assess Manag* 2020;00:1–9. © 2020 The Authors. *Integrated Environmental Assessment and Management* published by Wiley Periodicals, Inc. on behalf of Society of Environmental Toxicology & Chemistry (SETAC)

Keywords: Fish embryo toxicity   Acute fish toxicity   Weight of evidence   Bayesian network   Hazard assessment

INTRODUCTION

Many different chemical legislations (such as the European legislation for Registration, Evaluation, Authorisation and Restriction of Chemicals [REACH] Regulation [EC] 1907/2006 [EC 2006]) place an emphasis on the need for acute fish toxicity (AFT) data for the hazard assessment of chemicals. These data are required in combination with acute toxicity data from species of different trophic levels (i.e., algae and invertebrates) and are the minimum data requirements to perform an environmental risk assessment. However, the use of vertebrate organisms for acute ecotoxicity assessments has been challenged for ethical reasons and due to certain legislations (such as EU 2010) that specify that the use of vertebrate organisms should be avoided wherever possible. Consistent with the European Union (EU) Directive on the Use of Animals for Scientific Experimentation (EU 2010), REACH also specifies that the use of vertebrate organisms should be avoided wherever possible.

Russell and Burch (1959) first introduced the concept of humane experimental techniques for animal testing when they proposed the term "the 3Rs." These 3Rs refer to Reduction (in numbers of animals), Refinement (of any procedure), and Replacement (of vertebrate organisms). More

recently, the traditional 3Rs have been expanded to include 3 additional Rs, namely that any alternative should be Reliable/Robust and Relevant and gain Regulatory acceptance (OECD 1996; Lillicrap, Belanger et al. 2016). Henceforth, the traditional term of "3Rs" has since been refined and replaced to be considered as the "6Rs" of (eco)toxicity testing (Lillicrap, Belanger et al. 2016). For more than 20 y, scientists have been developing alternative approaches to predict AFT without the need to use fish. These include, but are not limited to, the use of quantitative structure–activity relationships (QSARs), in vitro test methods (e.g., the use of fish cells to determine cytotoxicity; Fischer et al. 2019), the use of other organisms such as invertebrates and algae (Rawlings et al. 2019), and the use of nonprotected life stages, in certain geographical regions, such as fish embryos (Busquet et al. 2014).

The fish embryo test (FET) (German DIN 2001; ISO 2007), was first considered as a promising alternative to the use of juvenile fish (48-h golden ide acute toxicity test) for assessing effluent toxicity by the German Federal Agency of the Environment. Subsequently, the method was submitted to the Organisation for Economic Co-operation and Development (OECD) as a new test guideline for the purposes of replacing the AFT test (OECD 1992). After significant international validation efforts (Busquet et al. 2014), an extension of the test from 48 to 96 h in duration, and omission of the term "replacement to fish acute toxicity" in the introduction of the test guideline (TG), the FET test was finally accepted as an OECD test guideline nearly 8 y later (OECD 2013). However, its universal acceptance as a replacement to the acute fish toxicity test (OECD 1992) has remained an issue because regulators, such as the European Chemicals Agency (ECHA), have not accepted it as a complete replacement (Sobanska et al. 2018). A conservative approach to the acceptance of the FET test has been argued due to the existence of some limitations (e.g., neurotoxic mode of action [MoA]) and/or remaining uncertainties (e.g., deviation of some narcotic substances) regarding the FET test (Sobanska et al. 2018). Furthermore, it was concluded that "the FET test alone is currently not sufficient to meet the essential information on AFT as required by the REACH regulation" (Sobanska et al. 2018). This is despite Sobanska et al. (2018) and other authors (Belanger et al. 2013) describing a near perfect 1:1 correlation between toxicity values (EC50 and LC50) from the FET and the AFT tests. Nonetheless, Sobanska et al. (2018) stated that the FET test "may be used within weight-of-evidence approaches together with other independent, relevant, and reliable sources of information." The use of weight of evidence (WoE) has been specified for REACH as an option to meet the information requirements of Annexes VII to X where "Animal tests can be avoided if there is a weight of evidence which points to the likely properties of a substance […] if there is sufficient information from several independent sources leading to the conclusion that a substance has (or has not) a particular dangerous property, while the information from each single source alone is regarded

insufficient to support this notion" (ECHA 2016). Understanding how this WoE may work in practice, to enable FET data to be used in place of AFT data, has been a major challenge. Therefore, we have previously developed a Bayesian network (BN) that incorporates FET data with multiple lines of evidence to provide a probabilistic estimate for AFT (Moe et al. 2020). In contrast to Moe et al. (2020), which describes both the model development and model training, the purpose of the present paper is to evaluate the performance of the BN for predicting AFT data from FET data along with other lines of evidence.

A BN is a probabilistic modeling methodology that is increasingly being used in ecological risk assessment (e.g., Landis et al. 2019) as well as more generally in environmental research (e.g., Barton et al. 2012; Moe et al. 2016). Bayesian approaches were used to integrate several biological lines of evidence to predict human skin sensitization to chemicals and is accompanied by a now elucidated adverse outcome pathway (OECD 2012b; Jaworska et al. 2015). A BN can integrate large amounts of data and other information sources by using discrete probability distributions and predicts the probability of specified states of selected variables. With regard to the FET–AFT BN, a specified state refers to, for example, a given interval of LC50 values for AFT. The purpose of the proposed BN model is to integrate information from large ecotoxicological and physicochemical data sets and apply it to predict AFT from data on FET tests, in combination with other relevant information in a WoE approach. The following sections describe the model setup and the evaluation of the model with a suite of different chemicals. It is envisaged that the use of the BN model will eventually fulfill the requirements of the regulatory community to accept FET data in place of AFT data.

MATERIALS AND METHODS

Chemical database selection

A database of ecotoxicity and other physicochemical properties for a range of different chemicals was obtained from Procter and Gamble. This database contained data for 237 substances and the following toxicity data:

- Algae: 264 EC50 values (duration 72 or 96 h, according to OECD 2006). The data comprised 7 algal species: the green algae *Chlorella pyrenoidosa*, *Chlorella vulgaris*, *Desmodesmus subspicatus*, and *Pseudokirchneriella subcapitata*; the diatom *Skeletonema costatum*; and the cyanobacteria *Anabaena flos-aquae* and *Microcystis aeruginosa*.
- Daphnids: 1164 EC50 values (48 h, according to OECD 2004). The 2 species used were *Daphnia magna* and *Daphnia pulex*.
- Juvenile fish: 1459 LC50 values (24, 48, 72, 91, or 96 h, according to OECD 1992). The data comprised 5 species: *Danio rerio* (zebrafish), *Lepomis macrochirus* (bluegill), *Oncorhynchus mykiss* (rainbow trout), *Oryzias latipes* (medaka), and *Pimephales promelas* (fathead minnow).
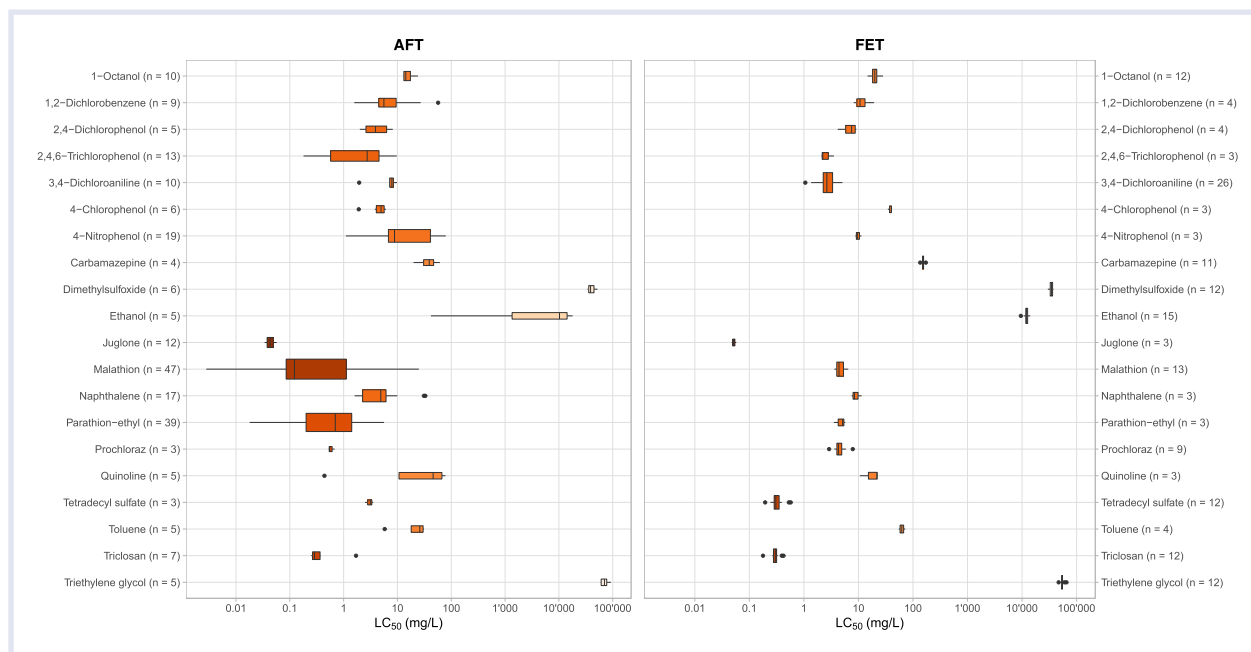
Figure 1. Overview of the data spread for the AFT data and the FET data used in the validation of the BN. Numbers in parenthesis, after each chemical name, are the number of data points (i.e., individual study data). AFT = acute fish toxicity; BN = Bayesian network; FET = fish embryo toxicity.

- Fish embryo: 541 LC50 values (48, 72, 96, 108, or 120 h, according to OECD 2013). Data were available for 4 species: *D. rerio*, *O. latipes*, *P. promelas*, and *Clarias gariepinus* (African sharptooth catfish).

The complete data set (i.e., data available for all 237 chemicals) was used for training the BN model, but a subset of chemicals was used to evaluate how accurately the model could predict AFT. The criteria for selecting these chemicals were that they had either a minimum of 3 FET and AFT data points or a minimum of 1 FET and AFT data point and a molecular weight >600 g/mol. This resulted in 28 candidate chemicals. An additional exclusion criterion was applied to remove any chemicals that had an extremely large spread of AFT data (e.g., for Cd the data varied by 5 orders of magnitude) or any chemicals that did not have any QSAR data. Even after the selection and exclusion criteria were applied, there were still some chemicals with quite a large spread of AFT data (see Figure 1). In comparison, the FET data were much less variable. The exclusion and selection criteria resulted in 20 prioritized chemicals (see Figure 1), which were subsequently used for the evaluation of the BN.
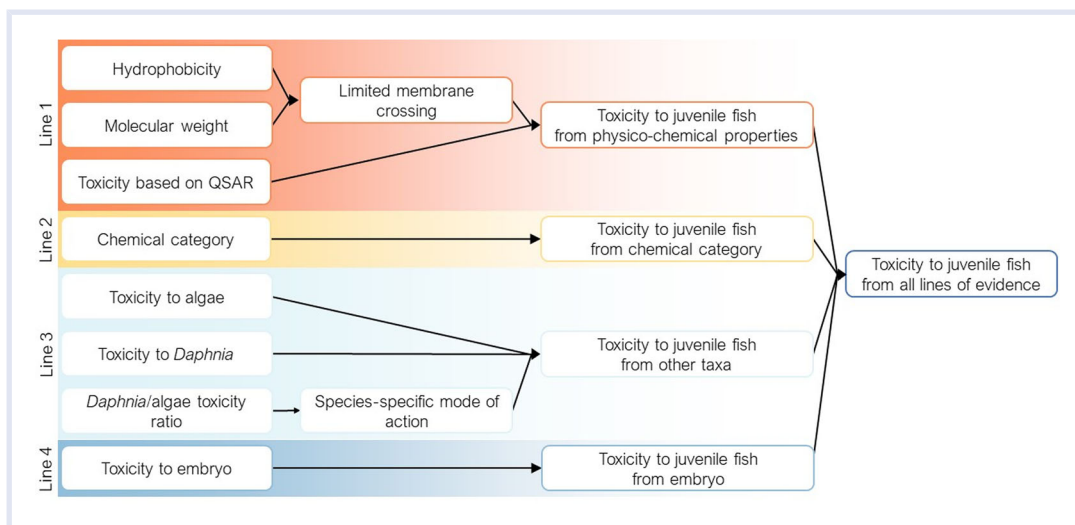
Description of the Bayesian network

The objective of this BN model is to predict the acute toxicity of a chemical to juvenile fish, corresponding to the interval of LC50 values from the AFT test, by integrating FET data with other relevant physicochemical and toxicological information. The application of this model is comparable to a WoE process as described by Suter et al. (2017) in which the assignment of prior and conditional probabilities to different variables corresponds to assigning weights to pieces of evidence. When predicting the AFT for a given

chemical, the calculation of posterior probability of each toxicity interval corresponds to the weighing of the total evidence for each hypothesis. The BN was implemented in the software HUGIN Researcher version 8.7, developed by HUGIN EXPERT A/S (http://www.hugin.com). The online web interface was made available through the demonstration web site (http://demo.hugin.com/FET). Parameterization and a description of the model construct is described in detail in Moe et al. (2020).

The BN model has 4 pathways (lines of evidence) for predicting the AFT of a given chemical (see Figure 2). These include data on 1) physical and chemical properties, 2) toxicity data to fish (AFT) from chemicals in the same category, 3) toxicity to other species (algae and daphnids), and 4) FET data. For the purposes of the present paper, it was important that all substances that were evaluated had a minimum of 3 values of corresponding FET and AFT data to enable suitable comparisons between the predicted AFT values and the observed AFT data.

Within each pathway (lines of evidence), discrete nodes (e.g., limited membrane crossing) were assigned. Each node has a conditional probability table (CPT) (e.g., see Table 1) which is conditional on the parent node (i.e., the prior input data point such as hydrophobicity). A full explanation of each of the CPTs for each node has been previously described (see Moe et al. 2020). The CPT values were obtained by 2 main methods: counts of observations, reflecting the distributions within the database, and expert judgment by the authors. All 4 pathways were assigned the same weight when combined in the predicted toxicity nodes. The toxicity intervals used in the BN model were discretized to 5 toxicity levels: very low (>100 mg/L), low (5–100 mg/L), medium (0.5–5 mg/L), high (0.5–0.01 mg/L), and very high (<0.01 mg/L).

Figure 2. Simplified illustration of the BN model with the 4 pathways (lines of evidence) contributing to predicting the AFT in a WoE approach. AFT = acute fish toxicity; BN = Bayesian network; WoE = weight of evidence.

It was recognized that variability within FET and AFT data would need to be accounted for in the BN. In Busquet et al. (2014), interlaboratory coefficients of variability (CVs) of the FET test were estimated to approximately 26% for all the substances evaluated in the international ring trial of the OECD TG 236 (OECD 2013). With regard to AFT, in an interlaboratory variability study (USEPA 2001), the CV was estimated to 20%. Therefore, as a conservative approach, the prior probabilities of all toxicity data in the input nodes were set at levels corresponding to minimum 30% CV. The BN predicts the toxicity level to juvenile fish for each chemical by combining information from all 4 pathways, and each of these pathways is described in the sections that follow.

Pathway 1, toxicity to juvenile fish based on physicochemical properties. Several descriptors of physicochemical properties can be used to estimate whether a chemical is likely to partition through biological membranes and hence increase the likelihood of cellular and/or molecular interactions with

Table 1. Example of conditional probability table (CPT) for the hydrophobicity and molecular weight nodes in pathway 1[a]

Hydrophobicity (log K_{ow})	Molecular weight (g/mol)	Probability for membrane crossing[b]		
		Low	Medium	High
<5.5	<600	0%	25%	75%
<5.5	>600	25%	50%	25%
>5.5	<600	25%	50%	25%
>5.5	>600	75%	25%	0%

[a] It illustrates the approach used for parameterization of the CPTs for the node "Limited membrane crossing": The probability of a substance crossing a biological membrane based on its physical properties (i.e., molecular weight and hydrophobicity).
[b] The probabilities are based on expert judgment.

the substance. One of the simplest metrics to determine whether the substance is bioavailable is the molecular weight (mw) or size of the molecule. These metrics have been included as one of the endpoints for persistent, bioaccumulative, and toxic (PBT) assessments within REACH, if used in a WoE approach (ECHA 2017). As indicated in Lillicrap, Springer et al. (2016), if the molecule has a physical size >4.3 nm and the molecule is unlikely to fold along linear structures (thus altering the length) or has a size (based on maximum diameter or Dmax$_{average}$) larger than 1.7 nm, then it is unlikely to pass across biological membranes. In addition, if the substance has a molecular weight above a certain size, then it should not be bioavailable, and according to ECHA (2017) substances with an average maximum diameter of >1.7 nm and a molecular weight of >1100 g/mol or >700 g/mol should not be bioaccumulative or very bioaccumulative, respectively (i.e., above these sizes they are increasingly nonbioavailable). However, these cutoff criteria should be considered with caution because not all molecules behave the same way, and according to Arnot et al. (2010) some of these assumptions may not have accounted for biotransformation of the substance occurring (i.e., providing a false positive assumption for bioavailability or not). For the purposes of this node in pathway 1, we have chosen a molecular weight cutoff from 600 g/mol to assume that substances with a molecular weight >600 g/mol are less likely to cross biological membranes. This is in line with Brooke et al. (1986), who indicated bioaccumulation potential had an upper limit of 600 g/mol.

In line with molecular weight, a measure of substance hydrophobicity can also be used as an indicator for assuming limited membrane-crossing potential. Hydrophobicity may be expressed by a substance's solubility in water being very low or based on the octanol–water partition coefficient (i.e., log K_{ow}) being very high. For the purpose of this second node of the first pathway, we have chosen hydrophobicity to be based on log K_{ow} and have

assigned a cutoff value of 5.5. This is in line with OECD TG 305 (OECD 2012a) in which substances with a log K_{ow} > 5.5 are recommended to be experimentally assessed using a dietary uptake exposure route, rather than an aqueous exposure route, due to reduced bioavailability. To account for the assumptions for potential membrane crossing, the conditional probability table (Table 1) in the BN model has 3 states: low probability of crossing (for large molecules), high probability (for small molecules), and medium probability to account for uncertainty in hydrophobicity or molecular size as a cutoff value (as detailed by Arnot et al. 2010). Other indicators, such as the Lipinski's rule of 5 (Lipinski et al. 1997), may be possible for accounting for limited uptake across biological membranes, but for the purposes of this BN, and to avoid too much complexity within the model, we have focused on only 2 molecular indicators in this pathway.

The third node of this first pathway is the use of QSAR data. The QSAR data are based on predicted values obtained from the United States Environmental Protection Agency (USEPA) Ecological Structure Activity Relationships (ECOSAR 1.11) Class Program (https://www.epa.gov/tsca-screening-tools/ecological-structure-activity-relationships-ecosar-predictive-model) and the Danish QSAR Database http://qsar.food.dtu.dk. The Danish QSAR database differs from the USEPA ECOSAR model because it has 2 different models (Leadscope and SciQSAR), which use different descriptors. The Leadscope model is based on structural features and numeric molecular descriptors (e.g., aLogP, polar surface area, number of H bond donors, Lipinski score, number of rotational bonds, parent atom count, parent molecular weight and number of H bond acceptors), and the SciQSAR model is based on molecular descriptors (e.g., molecular connectivity indices, molecular shape indices, topological indices, electrotopological (Atom E and HE-States) indices, and electrotopological bond types indices).

All data passed the preestablished criteria that was assigned as follows: For ECOSAR 1.11, class-specific QSAR models were used to predict AFT (96 h) if the equations passed set acceptability criteria ($R^2 \geq 0.6$, $n \geq 4$). If class-specific models were not available or did not meet these criteria, results from the Neutral Organic QSAR model were accepted. For the Danish QSAR database, extracted predicted values for Leadscope and SciQSAR fish 96-h LC50 values (*P. promelas*, fathead minnow) were used. Data were not used if the domain was listed as "OUT" in the Danish QSAR model. Solubility restrictions were applied and predicted values were excluded if they were 10 × the solubility values listed in the database. All QSAR fish LC50 results were than averaged for each chemical.

Pathway 2, toxicity to juvenile fish predicted by read across from chemical analogues with a similar MoA based on structural alerts. Of the extensive list of existing substances, there are already significant amounts of data that have been generated for determining the acute toxicity to fish. These chemicals can be categorized into functional groups that

may be considered similar, and it is possible to "read across" from 1 substance to another in the same category. Read across is already being used (and sometimes possibly misused) by registrants within REACH and other chemical legislations. However, it is quite reasonable to assume that extrapolating data from 1 chemical to another can either be logical or illogical, and for this reason, expert judgment is needed when applying read across. To avoid incorrect extrapolations being made, reliability frameworks (e.g., ECHA 2017) should be applied when assessing read across predictions.

The data used to develop this BN model included substances with different MoAs. Mode of action classes based on structural alerts were assigned by ECOSAR (v1.11), or by expert judgment if the chemical was outside of the domain. These included nonpolar narcosis, polar narcosis, uncoupler of oxidative phosphorylation, alkylation- or arylation-based reactivity, carbonyl reactivity (aldehyde), ester narcosis, organophosphate (OP)-mediated acetylcholinesterase (AChE) inhibition, hydrazine-based reactivity, pyridinium compounds, carbamate-mediated AChE inhibition, acrylate toxicity, diester toxicity, neurotoxicant: cyclodiene-type, neurotoxicant: pyrethroid, quinoline reactivity, and respiratory blocker. Where a chemical fitted a similar class of MoA based on structural alerts and corresponding AFT data were available, the AFT data were extracted and incorporated into the BN model (see Moe et al. 2020 for a full description of parameterization).

Pathway 3, toxicity to juvenile fish based on other species. The third pathway of the BN model incorporates ecotoxicity data from additional trophic levels. Environmental risk assessments incorporate hazard data from algae, daphnids, and fish. For that reason, we have included results from the algal inhibition assay (according to OECD [2006] TG 201) and the *D. magna* immobility assay (OECD [2004] TG 202). These trophic levels have been shown to be more sensitive than the AFT (Hutchinson et al. 2003; Jeram et al. 2005) and FET (Rawlings et al. 2019) between 75% to 80% of the time. Given that hazard classification and environmental risk assessment are based on the most sensitive taxa, algae and daphnids tend to routinely drive these assessments.

In pathway 3, we have assumed that the probability of high toxicity to fish is conditional on the chemical not having a species-specific MoA. For instance, if the chemical results in considerably different EC50 values for algae versus daphnids, this suggests that there is potentially a species-specific MoA affecting either algae or daphnids. To define what constitutes a difference, the ratio between the algae and invertebrate data is first calculated. Cutoff values for the ratio between the daphnid and algae data, as described in the European Centre for Ecotoxicology and Toxicology of Chemicals (ECETOC 2005), have been applied. For example, if the ratio of the 2 EC50 values is between 0.5 and 2, it can be assumed that these data are similar, and there is no specific MoA affecting either species. In these instances, it is reasonable that these EC50 values can be used toward

extrapolating an LC50 value for fish. Therefore, the CPT converts the EC50 values from algae and daphnids to LC50 values for fish with high precision (a narrow distribution). Conversely, a ratio >2 or <0.5 indicates that one of the species is more sensitive than the other, to a degree that suggests a species-specific MoA (e.g., herbicidal or insecticidal). In these cases, the probability that these EC50 data can be extrapolated to AFT LC50 data is low, and the CPT therefore has a wide distribution (high uncertainty). A full overview of the different conditional probabilities used, and a justification for each selection, is detailed in Moe et al. (2020), and an example of the prior probability distributions applied for both algae and daphnids is included in the Supplemental Data.

Pathway 4, FET based on experimental data. The fourth pathway is the use of FET data. These data need to be derived using the OECD TG 236 (OECD 2013) fish embryo acute toxicity test (FET). The data are then discretized into 5 levels:

very low (>100 mg/L), low (5–100 mg/L), medium (0.5–5 mg/L), high (0.5–0.01 mg/L), and very high (<0.01 mg/L).

Predicted toxicity to juvenile fish based on all 4 nodes. All 4 pathways of the BN model feed into the final node, which is an estimate of the predicted AFT based on molecular properties, fish data from other chemicals in the same category, ecotoxicity data from other species, and FET data. The final probability node is discretized into the same 5 categories for toxicity assignment as described in Pathway 4.

RESULTS AND DISCUSSIONS

The results from the BN predictions of the 20 prioritized chemicals are shown in Table 2. These results were compared to the actual measured AFT data to evaluate the accuracy of the predictions from the BN. The predicted toxicity interval for juvenile fish, based on the 4 different pathways, is also presented in Table 2. For the 20 chemicals evaluated

Table 2. Summary of Bayesian network model predictions for the 20 selected chemicals

Chemical	Chemical class (ECOSAR 1.11)	Verhaar classification	Observed juvenile[a]	Observed embryo[a]	BN prediction	Correct prediction[b]
1,2-Dichlorobenzene	Neutral organic	Nonpolar narcosis	Low	Low	Low	Y
1-Octanol	Neutral organic	Nonpolar narcosis	Low	Low	Low	Y
2,4,6-Trichlorophenol	Phenol	Polar narcosis	Medium	Medium	Medium	Y
2,4-Dichlorophenol	Phenol	Polar narcosis	Medium	Low	Low	N (D)
3,4-Dichloroaniline	Aniline	Polar narcosis	Low	Medium	Low	Y
4-Chlorophenol	Phenol	Polar narcosis	Medium	Low	Low	N (D)
4-Nitrophenol	Phenol	Polar narcosis	Low	Low	Low	Y
Carbamazepine	Substituted urea	Nonpolar narcosis	Low	Low	Low	Y
Dimethylsulfoxide	Neutral organic	Nonpolar narcosis	Very low	Very low	Very low	Y
Ethanol	Neutral organic	Nonpolar narcosis	Very low	Very low	Very low	Y
Juglone	Quinone	Alkylation/arylation based reactivity	High	High	High	Y
Malathion	Esters (dithiophosphates)	OP-mediated AChE inhibition	High	Medium	Medium	N (D)
Naphthalene	Neutral organic	Nonpolar narcosis	Medium	Low	Low	N (A)
Parathion-ethyl	Esters (monothiphosphates)	OP-mediated AChE inhibition	Medium	Low	Medium	Y
Prochloraz	Imidazole	Pyridinium compounds	Medium	Medium	Medium	Y
Quinoline	Neutral organic	Quinoline reactivity	Low	Low	Low	Y
Tetradecyl sulfate	Anionic surfactant	Nonpolar narcosis	Medium	High	Medium	Y
Toluene	Neutral organic	Nonpolar narcosis	Low	Low	Low	Y
Triclosan	Phenol	Polar narcosis	High	High	High	Y
Triethylene glycol	Neutral organic	Nonpolar narcosis	Very low	Very low	Very low	Y

A = algae; AChE = acetylcholinesterase; BN = Bayesian network; D = daphnid; OP = organophosphate.
[a] The observed juvenile and embryo are data obtained from the database.
[b] For the chemicals for which there was an incorrect prediction, the most sensitive species is identified in parenthesis.

herein, the BN model correctly predicted the toxicity interval for 80% of the substances. An example of the BN model, showing the distributions for each node for the chemical tetradecyl sulfate (chemical category anionic surfactant), is shown in the Supplemental Data. The AFT data obtained from the database (based on 3 observations) for tetradecyl sulfate indicated that it would be classified as causing medium toxicity (i.e., LC50 0.5–5 mg/L) to juvenile fish. The node "Toxicity to embryo (level)" was almost 100% high toxicity (i.e., 0.5–0.01 mg/L), whereas the predictions from the other lines of evidence were centered around low-to-medium toxicity. The resulting predicted toxicity to juvenile fish had the highest probability (41%) of the medium state (i.e., LC50 0.5–5 mg/L), which was consistent with the measured AFT data. In this case, using FET data alone would have overestimated the risk to juvenile fish, whereas FET data, in combination with the other lines of evidence, resulted in a more accurate prediction on toxicity to juvenile fish.

For the remaining 20% of the substances, where there was an incorrect prediction, either the daphnid or algae data were more sensitive in all cases. For the 3 substances, 2,4-dichlorophenol, 4-chlorophenol, and malathion, daphnids were most sensitive. These compounds are classified, according to the Verhaar classification system, as having a polar narcosis MoA for the 2 phenols, and OP-mediated AChE inhibition for malathion (Verhaar et al. 1992). Given that malathion is an insecticide, it is not surprising that daphnids were the most sensitive species. For the remaining substance naphthalene, which is a neutral organic causing nonpolar narcosis, algal growth inhibition was the most sensitive endpoint. For these 4 chemicals, where there was an incorrect prediction of the AFT, either daphnids or algae would have driven the environmental classification.

It is important to note that it was not possible, within the present study, to consult each study record for each data point to perform a reliability evaluation. Therefore, the comparisons between the BN predictions and the AFT data herein assume that all data were reliable. However, it is possible that some data may not be reliable, or that the data could be questionable. As an example, AFT data generated from different laboratories, and from different species of fish, can vary by multiple orders of magnitude (Belanger et al. 2013). In the initial exclusion exercise to identify the prioritized chemicals used to validate the BN, the AFT data for some substances, such as Cd, varied by more than 5 orders of magnitude and were excluded from the evaluations. The reason for such high variability may be due to the applicability of the test, differences between species sensitivities, or more likely, poor experimental data. It may also be due to other confounding factors such as when AFT data are based on nominal versus measured concentrations. To illustrate this, Jonker et al. (2018) performed an international ring trial on passive sampling and attributed the large interlaboratory variability that was observed to analytical chemistry. Only when 1 single laboratory performed the analytical chemistry for the ring trial was it possible to eliminate the large source of variability.

Furthermore, the appropriateness of the AFT test design (OECD 1992) may also be questionable. For instance, for hydrophobic substances (i.e., with a log K_{ow} > 5.5) a 96-h duration may not be sufficiently long enough to ensure that equilibrium within the fish and the exposure media has been achieved. This means that for certain hydrophobic substances, the AFT test may be underestimating the actual toxicity because a critical body burden has not been achieved within the fish. Similarly, certain embryo toxic substances, such as triazoles and glycol ethers, which cause growth retardation and malformations in zebrafish embryos (Hermsen et al. 2011), will not elicit the same response in juvenile fish. Henceforth, given that the AFT is a relatively crude test (i.e., do the fish live or die), or significantly underestimates the toxicity due to the exposure duration being too short, or there is a species-specific MoA (e.g., herbicide or insecticide), the other baseline toxicity assays (e.g., algae and invertebrates) are often more sensitive than fish. To illustrate this point, where the BN did not correctly predict the AFT (20% of the chemicals), it was inconsequential because algae or daphnids were more sensitive in all cases. At this stage, it should be noted that the FET test is also not without its limitations. For example, it has been shown that the FET test exhibits a weak response to substances that have a neurotoxic MoA (Klüver et al. 2015). However, Klüver et al. (2015) recommended that substances with a neurotoxic MoA could be identified using behavioral endpoints such as embryonic locomotion. In the future, it might be possible that more sensitive endpoints, such as behavior, may or should be incorporated into the OECD TG 236 (OECD 2013) to account for specific MoA such as neurotoxicity.

Clearly, data from only 1 ecotoxicity test (e.g., algal toxicity, daphnid immobilization, or AFT) are insufficient to provide adequate information to perform a hazard and risk assessment or for the purposes of chemical classification. Therefore, it is imperative that all environmental data be incorporated to provide better confidence in performing environmental risk assessments and for classification purposes. To this end, the requirement to develop a WoE approach has been welcomed by the authors of the present paper because we believe that our BN model is timely and of greater importance than to simply predict AFT data from FET data. Moreover, the use of our BN model will help to improve future environmental risk assessments of chemicals per se. However, the model that has been developed thus far is based on currently regulatory accepted methods and approaches, and there are many other lines of evidence that could be incorporated to increase the predictive power of the model. One example would be the use of cytotoxicity data from the RT-gill cytotoxicity assay, which has recently been accepted as an international standard (ISO 2019). Another line of evidence would be to incorporate information related to metabolism and neurotoxicity to inform those substances that might require metabolic activation, or that have a specific MoA that causes toxicity to older life stages (e.g., juvenile) of fish. Furthermore, data from other

sources of information (e.g., other species, chronic toxicity data, or human safety information) or other QSAR models could also refine the current model. With these additional pieces of information, it is envisaged that the BN model could improve the WoE to more accurately predict AFT data, enabling FET data to be submitted in place of AFT data for regulatory requirements such as REACH.

CONCLUSIONS

A BN has been developed and is able to predict AFT from FET data, in combination with other lines of evidence. For a subset of chemicals, the BN was able to accurately predict the toxicity interval for 80% of the chemicals evaluated. In these cases, the BN demonstrated that a sufficient WoE could justify the use of FET data in place of AFT data. For the remaining 20% of the chemicals, where an incorrect prediction was made, either the daphnid or algae data were more sensitive, and these data would have driven any environmental classification. Nonetheless, the current BN model could benefit from additional lines of evidence to be included, and a larger database of chemicals to further train the model to reduce any uncertainties in the predictions. It is recommended that BN models should be used more often for determining WoE, and we encourage further dialogue with the scientific and regulatory community to advance the acceptance of such models to replace the use of AFT tests in the future.

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Data Availability Statement—Data in the database are available on the Wiley Online Library at https://doi.org/10.1002/etc.4351. All other data can be made available on request.

SUPPLEMENTAL DATA

Figure SI-1. Prior probability distributions applied in the BN for the nodes [algae node name] and [daphnids node name]. These prior probability distributions are derived from the training set and correspond to the frequency distributions of toxicity values in the given concentration intervals. BN = Bayesian network.

Figure SI-2. Example of the BN model, showing the different distributions for each node for the chemical tetradecyl sulfate. BN = Bayesian network.

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