

Web-based InterspeciesCorrelation Estimation (Web-ICE)for Acute Toxicity

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12 February 2014



http://www.epa.gov/ceampubl/fchain/webice/

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What is Web-ICE?

An internet application developed by the US
 Environmental Protection Agency, Office of Research and Development, Gulf Ecology Division.



- Uses interspecies correlation estimation (ICE) models to estimate acute toxicity to a species, genus, or family from the known toxicity of a surrogate species.
- Contains modules to derive acute hazard levels (e.g., HC5) and endangered species toxicity useful to chemical hazard assessment and Ecological Risk Assessment (ERA).

How can Web-ICE valuable to ERA?

Estimates acute toxicity (LC/EC50) for a species, genus or family from a surrogate species

ICE in ERA

- Populates toxicity database
- Direct toxicity estimation for endangered species
- Allows for species sensitivity comparisons and development of SSDs







Atlantic salmon

Discussion Outline

1) Technical Basis of ICE models

- Database & model development
- Validation and uncertainty analysis
- User guidelines / Rules of Thumb



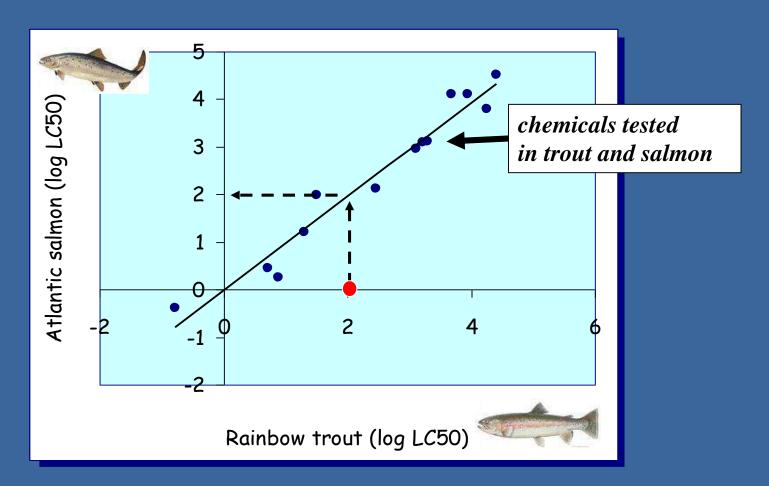
2) Applications in SSDs

10:30 – 12:00 Web-ICE Demonstrations

Interspecies Correlation Estimation (ICE)

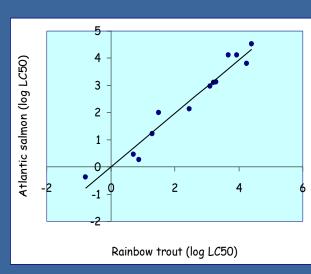
Log-linear least squares regressions of the relationship between the acute toxicity (LC50/LD50) of chemicals tested in two species.

 $Log(Toxicity_{predicted}) = a + b*Log(Toxicity_{surrogate})$



ICE Models – the basics

- 1. ICE models start with large database of acute toxicity
 - e.g. ICE v3.1 aquatic invert/fish database = 5487 LC50 values; 1258 chemicals; 180 species
- 2. All possible pairings of species by common chemical
 - e.g. 180*180 = 32400 potential pairings
- 3. ICE model = Log-linear least squares regression of common chemicals tested in two species
 - some pairings will not yield any ICE model
 - some models will not be significant (p>0.05)
- 4. Suite of ICE models dependent on toxicity database



Web-ICE Databases

Aquatic (fish and invertebrates):

- US EPA Office of Pesticide Programs Ecotoxicity Database
- US EPA Office of Pollution Prevention & Toxics PMN, HPV
- US/EPA/ Office of Research and Development
 - Ecotox
 - e.g. Mayer 1987
- Ambient Water Quality Criteria
- US Geologic Survey
- Mayer and Ellersieck 1986
- Open literature sources
 - endangered species
 - mollusks



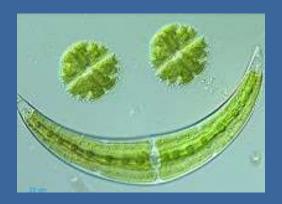


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Web-ICE Databases

NEW Aquatic (algae):

- Procter & Gamble (CRADA)
- EPA Office of Pesticide Program
- Open literature searches
- Ecotox



Wildlife (birds and mammals):

- US EPA Office of Pesticide Programs Ecotoxicity Database
- Environment Canada (Baril et al. 1994)
- Hudson et al. 1984
- Schafer et al. 1983
- Shafer and Bowles 1985
- Safer and Bowles 2004
- Smith 1987



Data Standardization

All Databases

- cas/name consistency
- single compound tested
- a.i. > 90%
- chemical & element specific AWQC normalizations (e.g. ammonia, metals)

Aquatic animal

- no sediment, dietary, mixed dose exposures or phototoxicity results
- ASTM/OPPTS standards
- endpoint = mortality/immobilization
- 48h EC/LC50 daphnids, midges and mosquitoes
- 96h EC/LC50 fish and other invertebrates
- juvenile only: fish, amphibians, insects, molluscs, decapods
- other groups: all life stages

Aquatic plants (algae)

• 72 or 96 hour EC50, growth rate or biomass

Wildlife

Single oral dose LD50 (mg/kg), adults only

Web-ICE Databases

Taxa	# Records	# Chemicals	# Species
Aquatic animal	5487	1258	180
Algae	1647	457	57
Wildlife	4329	951	156

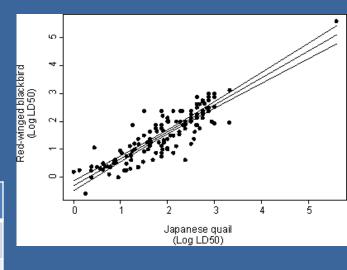


Model Development

Models are only developed for species within the same database (e.g., no algae to fish models)

Log10(predicted toxicity) = a + b*Log10(surrogate toxicity)

Taxa	Species	Genus	Family
Aquatic animal	780	289	374
Algae	58	44	0
Wildlife	560	0	292



- All species/taxa are paired with each other by common chemical within single database
- Geometric means for more than one toxicity value per species/chemical pair
- Excluded species/chemical pairs where the minimum and maximum tox values were greater than 10-fold
- Three or more common chemicals per pair are required to develop a model
- Only models with a significant relationship (p-value < 0.05) are included in Web-ICE

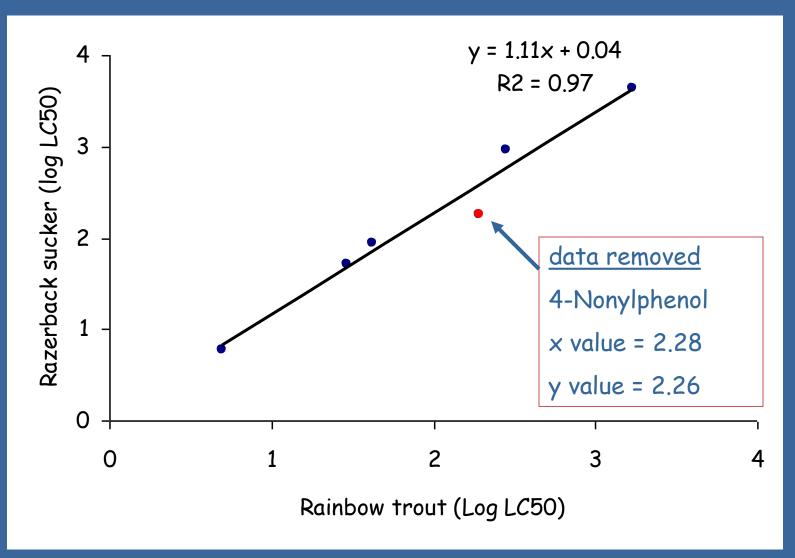
Model Validation & Uncertainty Analysis

- Only included models significant at the p<0.05 level
- Leave-1-out cross-validation performed on models N ≥ 4

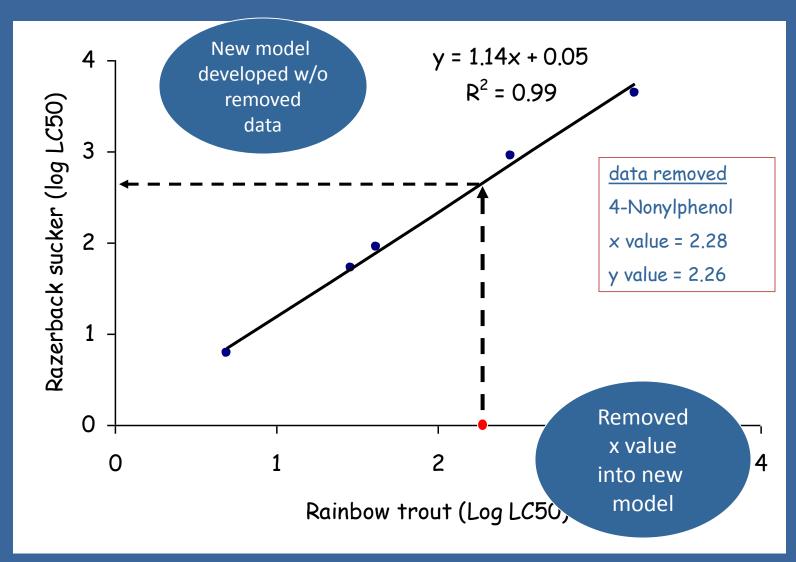
leave-1-out cross-validation

Each data point is removed, one at a time, and the model is rebuilt with remaining data. Removed surrogate data are used to estimate removed predicted data from rebuilt model.

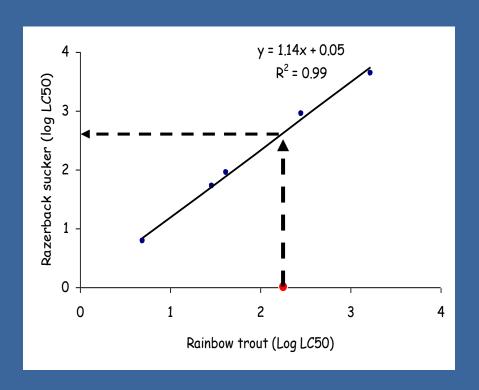
Model Validation: Leave-1-Out



Model Validation: Leave-1-Out



Model Validation: Leave-1-Out



data removed

4-Nonylphenol

x value = 2.28

y value = 2.26

New model predicts 2.64

compare to actual value (2.26)

"fold difference" of untransformed data=

 $10^{(2.64)}/10^{(2.26)} = 2.46$

*** "fold diff" is maximum of predicted/actual or actual/predicted

Model Uncertainty Analyses

- Used "fold difference" of cross-validation to identify areas of model uncertainty
- Species level models
 - Aquatic: 10914 data points removed and validated from 696 models
 - Wildlife: 11846 data points from 538 models
- Model "Cross-validation success rate" = % removed data within 5-fold
- Taxonomic Distance
- Model parameters
- Mode of Action

Taxonomic Distance

Measures the taxonomic relatedness of the surrogate and predicted taxa

Surrogate:
Rainbow trout



Taxonomic distance

shared taxonomic level

1 – genus

Cutthroat trout (Oncorynchus clarki)



2 - family

Atlantic salmon (Salmo salar)



3 - order

4 - class

Sheepshead minnow (Cyprinodon variegatus)



5 - phylum

S. leopard frog (Rana sphenocephala)



6 - kingdom

Daphnid (Daphnia magna)



Model uncertainty & taxonomic distance

Aquatic animals in same family ~ 90% within 5-fold, 95% within 10-fold

Shared	Significant _ models (N)	Percentage within predicted range			
taxonomic level		5-fold	10-fold	50-fold	> 50-fold
Genus (1)	254	94	2	3	1
Family (2)	700	91	5	3	1
Order (3)	208	86	10	3	0
Class (4)	4432	80	9	8	3
Phylum (5)	764	61	14	17	8
Kingdom (6)	4556	56	15	19	10

Model uncertainty & taxonomic distance

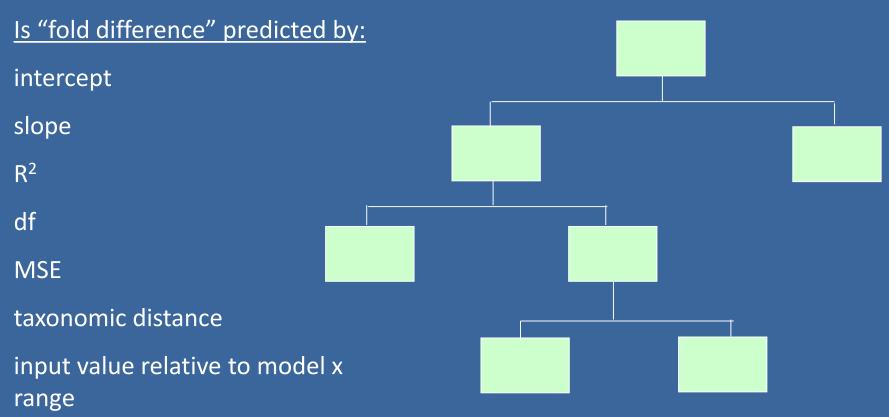
Wildlife in same order ~ 90% within 5-fold, 97% within 10-fold

Percentage of all datapoints in cross-validation category

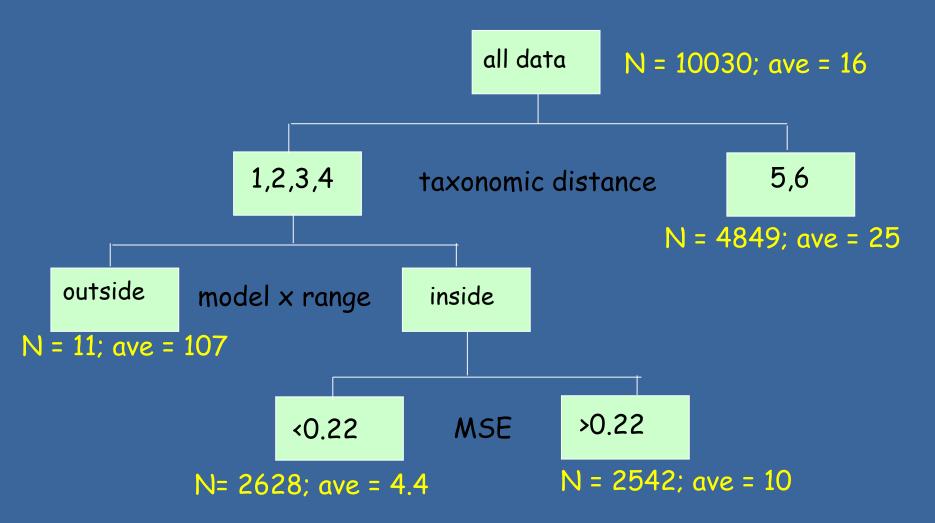
Common level	Number datapoints	5-fold	10-fold	50-fold	> 50 fold
Genus (1)	48	100	0	0	0
Family (2)	1452	92	6	2	0
Order (3)	2238	90	7	3	0.3
Class (4)	5706	85	10	5	0.2
Phylum (5)	2402	76	13	9	1.5

Uncertainty and Model Parameters *Aquatic*

Regression Tree:



Regression Tree - Aquatic (Decision Tree, Pattern Recognition)



User Guidance: Selecting a Model with Low Uncertainty

No one attribute defines model robustness!

Rules of Thumb:

- low MSE (< 0.94) (based on re-analysis)
- $high R^2 (> ^0.6)$
- Close confidence intervals (input value should be relatively close to the range of surrogate data used to create the model)

Weight of Evidence

- high cross-validation success rate
- close taxonomic distance (within class)





Applications: Species Sensitivity Distributions

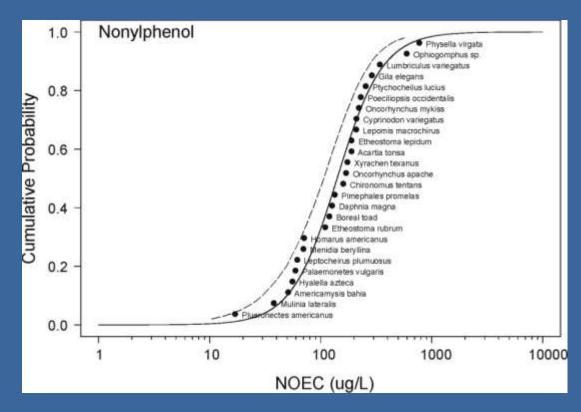
• Supplement datasets to populate SSDs

Develop SSDs from single toxicity value

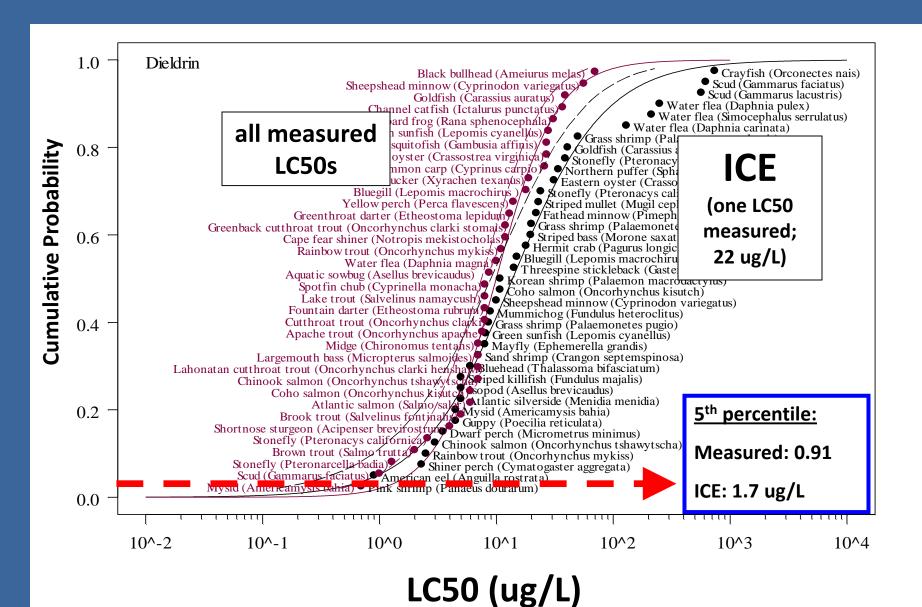
link to QSAR

ICE-generated SSDs are comparable to SSDs

built from measured data



ICE and Measured SSDs



How Do ICE and Measured SSDs Compare for Aquatic Organisms?

- Aquatic Species Proof of Concept (Dyer et al. 2006)
 - ICE-based SSDs had HC5s within an order of magnitude of measured HC5s
- Aquatic Species, Expanded Study (Dyer et al. 2008)
 - 55 AWQC chemicals, 4 surrogate species (P. promelas, O. mykiss, C. varigatus, D. magna)
 - Using fish surrogate to predict fish and invertebrate surrogate to predict invertebrates yielded HC5s similar to measured (average factor of 3.0 ± 6.7 over 7 orders of toxic magnitude)

Environ. Sci. Technol. 2008, 42, 3076-3083

Comparison of Species Sensitivity Distributions Derived from Interspecies Correlation Models to Distributions used to Derive Water Quality Criteria

SCOTT D. DYER,***
DONALD J. VERSTEEG,*
SCOTT E. BELANGER,* JOEL G. CHANEY,*
SANDY RAIMONDO,* AND
MACE G. BARRON*

The Procter and Gamble Company, 11810 East Miami River

Introduction

Ecological risk assessments typically require characterizing the effects of multiple chemicals on a diversity of ecological receptors using toxicity data for only a limited number of species. Regulatory activities such as REACH (Registration, Evaluation and Authorization of Chemicals (1, 2); http://europa.eu.int/comm/environment/chemicals/reach.htm), ICCA (International Council of Chemical Associations) High Production Volume (HPV) Chemicals Challenge (http://www.icca-chem.org/), and Canada's Domestic Substance List (3) will also create new demands for toxicity data. Note that the ICCA HPV Chemicals challenge is in cooperation with OECD and its member countries which then also include other HPV challenge programs such as those of the United States Environmental Protection Agency (USEPA) and Japan.

Augmenting Datasets with ICE-Generated Data

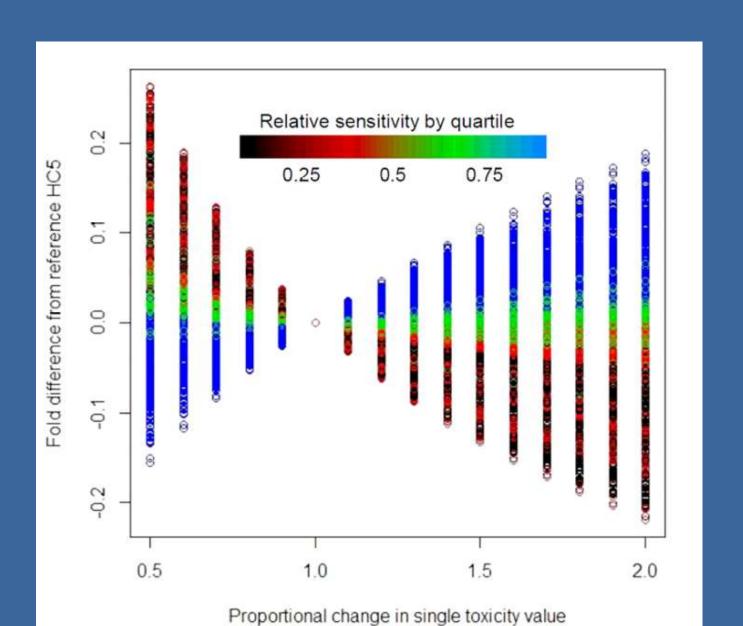
- Limited measured data augmented with ICE toxicity values (49 chemicals with > 10 species)
- SSD sensitivity analysis
- ICE prediction uncertainty contributed less to HC5 variability than species composition
- ICE prediction variability influence on HC5 dependent on location in SSD

Hazard/Risk Assessment Environmental Toxicology and Chemistry
DOI 10.1002/etc.2456

AUGMENTING AQUATIC SPECIES SENSITIVITY DISTRIBUTIONS WITH
INTERSPECIES TOXICITY ESTIMATION MODELS

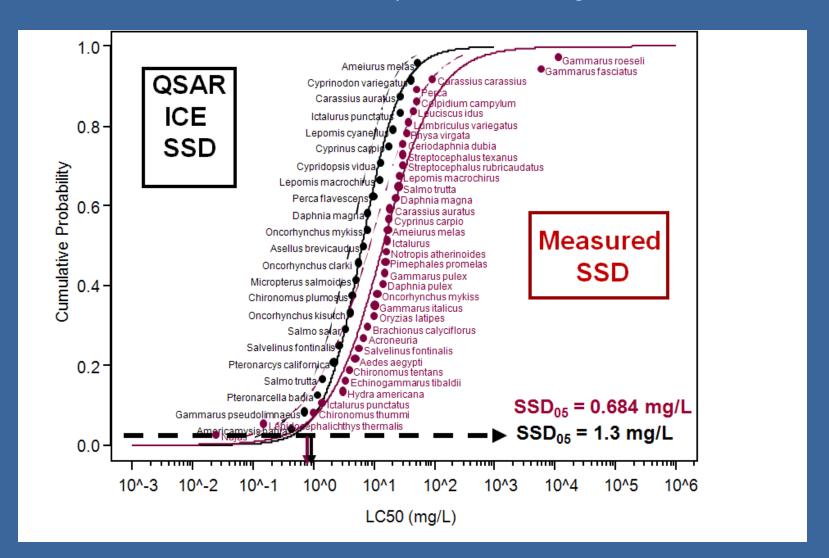
JILL A. AWKERMAN*, SANDY RAIMONDO, CRYSTAL R. JACKSON, and MACE G. BARRON

Influence of SSD data variability on HC5



Web-ICE and QSAR

QSAR tox estimate used as input into ICE SSD generator



QSAR → Web-ICE → SSD

- in silico approach
- 10 chems/4 MOAs; daphnid, fathead
- Compared measured SSD HC5 to in silico HC5
 - Results variability with MOA & QSAR uncertainty
 - Variability reduced with identical species composition
 - 80% HC5s within 5-fold of measured using ECOSAR QSAR value



Evaluation of in silico development of aquatic toxicity species sensitivity distributions

Mace G. Barron*, Crystal R. Jackson, Jill A. Awkerman

U.S. EPA, GED, 1 Sahine Island Drive, Gulf Breeze, FL 32561, USA

How Do ICE and Measured SSDs Compare for Wildlife?

- Used ICE database of acute toxicity (23 chemicals with > 15 species)
- For each chemical, removed records from database to generate independent ICE models
- Created SSDs
 - 1. measured data
 - 2. ICE data 5 surrogates (Japanese Quail, Mallard, Northern bobwhite, redwinged blackbird, Norway rat)
- 100% HC5s within a factor of 5 using bird & mammal surrogates

Environ. Sci. Technol. 2008, 42, 3447-3452

Development of Species Sensitivity Distributions for Wildlife using Interspecies Toxicity Correlation Models

JILL A. AWKERMAN, SANDY RAIMONDO, AND MACE G. BARRON

U.S. Environmental Protection Agency, Gulf Ecology Division, 1 Sabine Island Drive, Gulf Breeze, Florida 32561 species with known uncertainty (2). In practice, a leastsquares model II regression is developed from the known
LD50 values of two species tested for the same group of
chemicals. The model is then used to approximate toxicity
of one species to a chemical from the known toxicity of the
other species (the surrogate). ICE models for wildlife have
been determined to be most accurate for two species within
the same order, but may be used to estimate toxicity from
birds to mammals (2). Confidence intervals calculated with
ICE estimates are based on the linear relationship and mean
model error and may be used to assess the robustness of
model predictions. In aquatic species, acute toxicity values
generated from ICE models have been used to populate SSDs
and have been recommended for generating reasonable

Can we use only rodent data and ICE to develop reliable wildlife SSDs?

- Database of rodent oral dose LD50 data (Hazardous Substances Data Bank)
- Mouse: 109 chemicals; Rat: 175 chemicals
 - Rodents most often in least sensitive quartile; not great surrogates
- Created measured and surrogate ICE SSDs:
 - rat (53 chems) and mouse (32 chems)
- Compared HD5s
 - 78% of HD5 values within 5x of measured

Journal of Textenings and Environmental Health, Part.A, 72: 1604–1609, 2008 Copyright © Taylor & Friencis Group, LLC ISSN: 1528-7394 print / 1087-2620 online DOI: 10.1080/15287390003333401



Estimation of Wildlife Hazard Levels Using Interspecies Correlation Models and Standard Laboratory Rodent Toxicity Data

Application of Web-ICE

Defensibility

- cross-validation of models provides user with estimate of model performance
- all models significant



User guidance

- clearly outlined
- uncertainty analyses identify "Rules of Thumb" (Aquatic complete in 2008)
- Web-ICE alerts users to potentially ill-fitting data

Demonstration of use in SSDs

- Aquatic and wildlife
- Species composition
- Taxonomic distance of surrogate and predicted





http://www.epa.gov/ceampubl/fchain/webice/