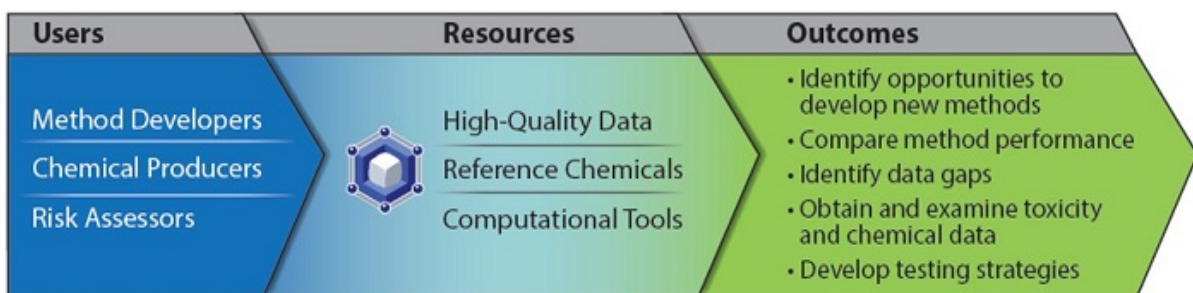


User-friendly Toxicology Tools from ICE Workflows

S. Bell¹, J. Phillips², N. Cariello¹, P. Ceger¹, X. Chang¹, F. Hermes¹, K. Mansouri¹, R. Rai¹, D. Allen¹, W. Casey³, N. Kleinstreuer³

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The Integrated Chemical Environment



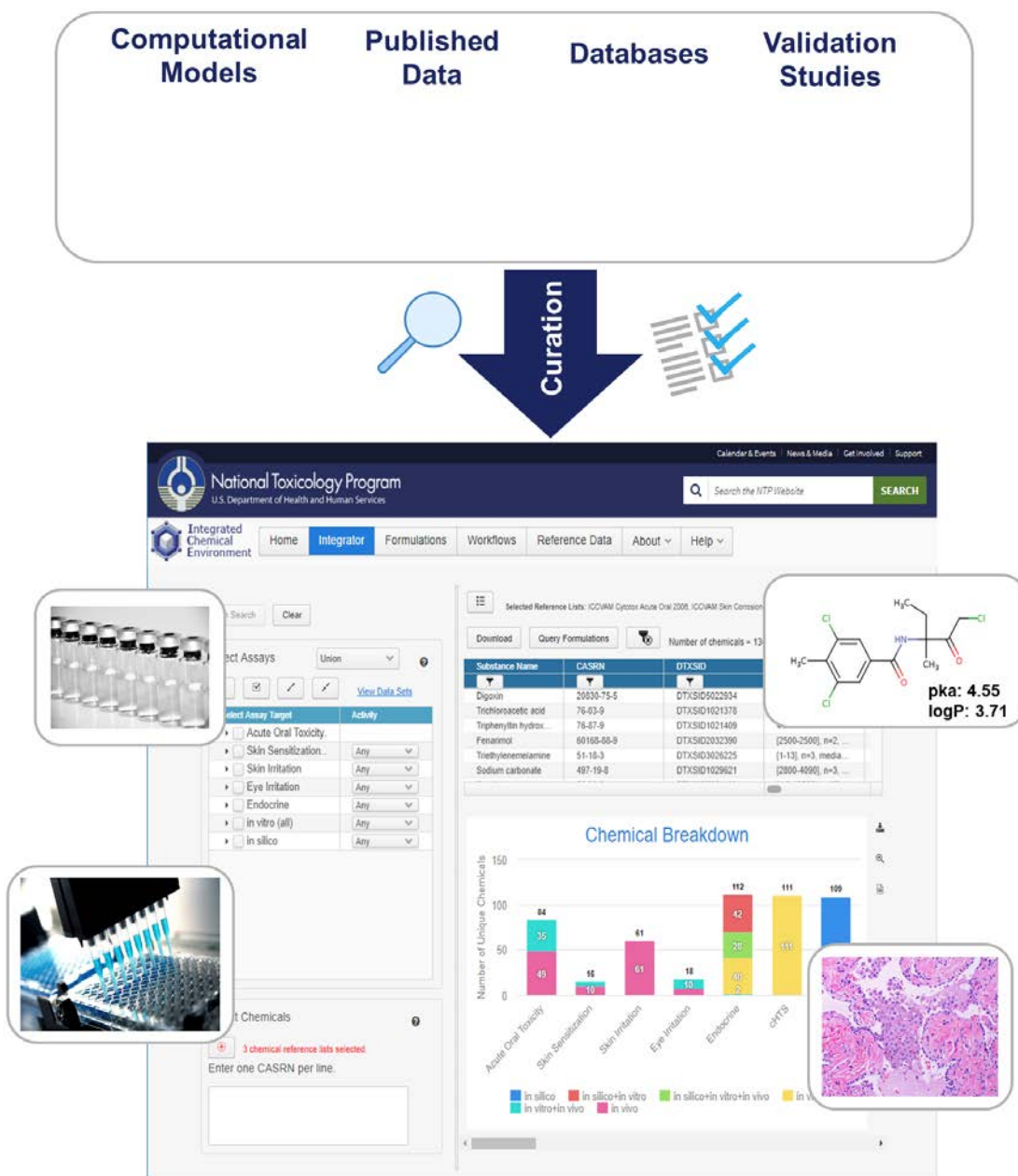
ICE provides free online access to:

- Curated in vivo and in vitro test data
- In silico toxicity predictions and chemical property data
- Reference chemical lists
- Computational tools and workflows related to chemical characterization and predicting toxicity

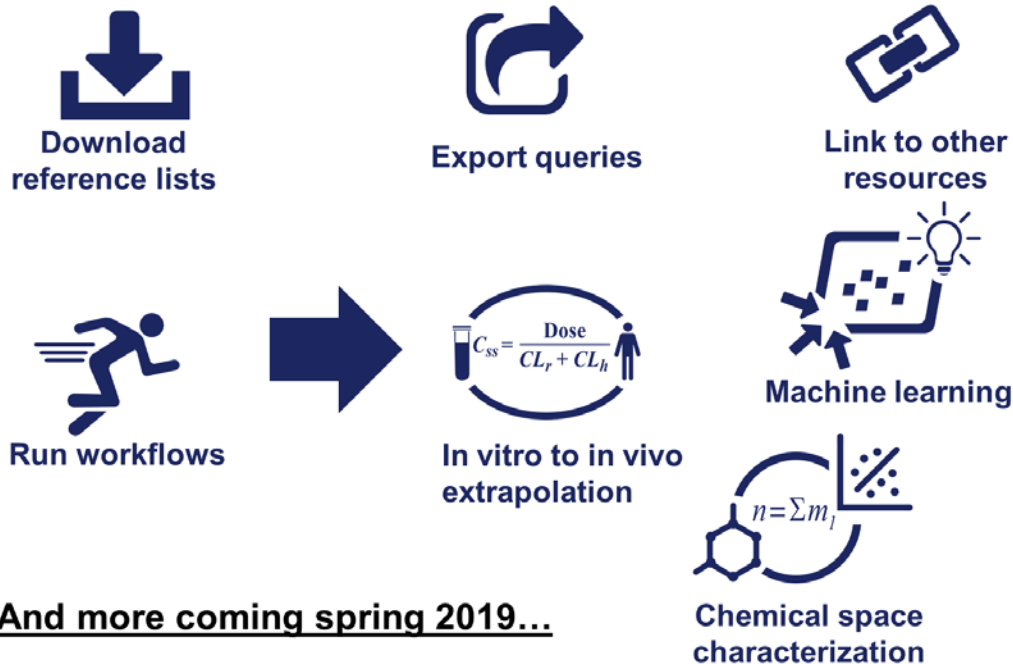
ICE supports:

- Data integration: brings together available data, including data on formulations
- Results exploration: enables dynamic, graphical exploration with publication-quality graphics
- Data analysis: allows characterization of data using online workflows
- FAIR data access

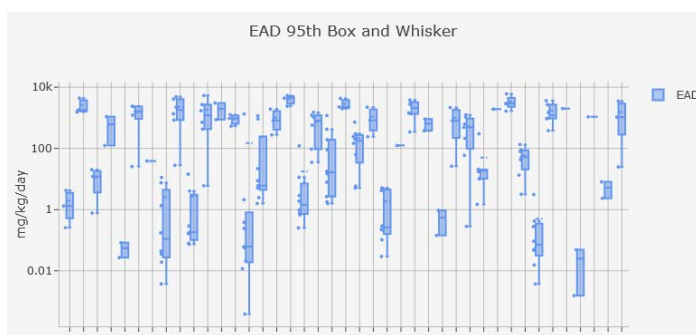
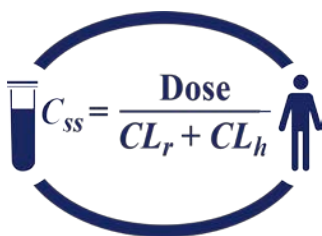
What Goes into ICE?



Making Data and Tools Available



In Vitro to In Vivo Extrapolation



Use high-throughput in vitro data available from ICE to estimate external dose via a one-compartment model

IVIVE tools planned for future releases:

- Three-compartment physiologically based pharmacokinetic model
- Glucuronidation and sulfation predictions

- Reverse toxicokinetics predictions
- Calculations with user-provided data on ICE platform

Machine Learning



Use the machine learning tool for hypothesis generation and to explore different machine learning approaches using ICE data. Selected endpoints (for classification and regression modeling) and algorithms are available through ICE to facilitate the ease of use by those with limited background in computational toxicology

Endpoints for prediction

- LLNA: call
- Uterotrophic: call, LEL
- Human Potency: call, NOEL/LOEL

Machine learning methods available

- rf: random forest
- rpart: recursive partitioning
- knn: k-nearest neighbor
- svmRadial: support vector machine with a radial kernel

- pls: partial least squares regression

Models: knn, rf
Metrics: ROC, Sens, Spec
Number of differences: 1
p-value adjustment: bonferroni

[1] ----- knn -----
[1] Machine Learning
Confusion Matrix and
Reference
Prediction Active :
Active 61
Inactive 4

Accu : 0.9564
95% CI : (0.8914, 0.9842)
No Information Rate : 0.625
P-Value [Acc > NIR] : 4.602e-15

Kappa : 0.899
McNemar's Test P-Value : 0.3711

Sensitivity : 0.9385
Specificity : 0.9744
Pos Pred Value : 0.9839
Neg Pred Value : 0.9048
Prevalence : 0.6250
Detection Rate : 0.5865
Detection Prevalence : 0.5962
Balanced Accuracy : 0.9564

'Positive' Class : Active

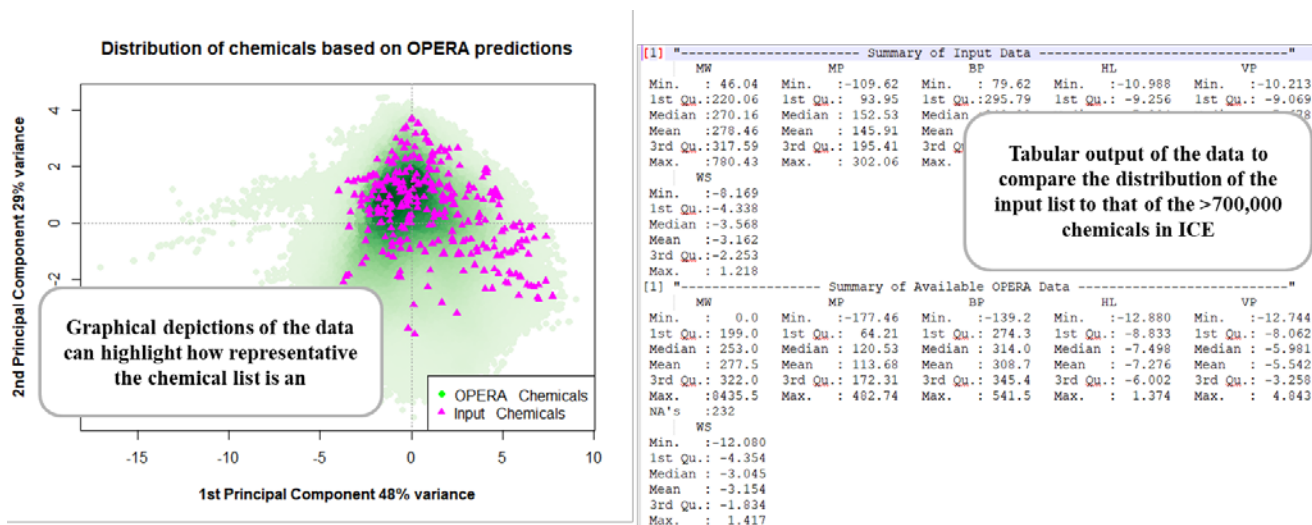
	A	B	C	D	E
1 CASRN	knn	svmRadial	toPredict	PhysChem	
2 95-70-5	Active	Active	Active	0.355	
3 26172-55-	Active	Active	Active	0.355	
4 81406-37-	Active	Active	Active	0.355	
5 119-36-8	Inactive	Inactive	Inactive	0.355	
6 87-86-5	Active	Active	Active	0.355	
7 862892-90	Active	Active	Active	0.355	
8 531-95-3	Active	Active	Inactive	0.355	
9 66422-95-	Active	Active	Active	-0.816	
10 122-57-6	Active	Active	Active	1.826	
11 150114-71	Inactive	Inactive	Inactive	0.675	
12 1154-59-2	Active	Active	Active	4.99	
13 89-25-8	Active	Active	Active	0.929	

Performance statistics of the model including the confusion matrix (classification) and RMSE (regression) are available to compare method performance

Machine learning tool outputs a table with the predictions from each model along with the data used for training the model

Chemical Characterization

Leverage ICE models to characterize a chemical list, getting information on the chemical space covered (or not covered) based on different physchem properties.



In the pipeline:

- Generate physchem and other structure-based predictions from user-provided chemical lists
- Prediction of chemical parameters for use in modeling (example. fraction unbound, pka)
- Chemical use category overview provided by EPA's Consumer Products Database (cpDAT)
- Integration with ChemMaps (<http://www.chemmaps.com/>)

Contact Us



To get announcements of ICE updates and other NICEATM activities, visit the NIH mailing list page for NICEATM News at <https://list.nih.gov/cgi-bin/wa.exe?SUBED1=niceatm-l&A=1> and click “Subscribe.”

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