



A Comprehensive Primer for the CompTox Chemicals Dashboard

- **Antony Williams** – General chemistry capabilities and advanced search, predicted model and experimental data, real time modeling, advanced searches based on mass, formula, formula generation (15 minutes)
- **Katie Paul Friedman** – Hazard and Bioactivity Functionality (15 minutes)
- **John Wambaugh** – Absorption, Distribution, Metabolism and Excretion Data (10 minutes)
- **Kathie Dionisio** – Exposure Functionality including Product Use Categories, Chemical Weight Fraction and Chemical Functional Use (10 minutes)
- **Kristin Isaacs** – Exposure Functionality including Toxics Release Inventory, Monitoring Data, Production Volume and Exposure Predictions (10 minutes)
- **Grace Patlewicz** – GenRA, Similar Compounds and Related Substances (10 minutes)
- **Nancy Baker** – Literature Search Functionality (15 minutes)
- **Antony Williams** – Batch search and lists (10 mins)



A Comprehensive Primer for the CompTox Chemicals Dashboard

**General chemistry capabilities and advanced search,
predicted model and experimental data, real time
modeling, advanced searches based on mass, formula,
formula generation**

Antony Williams

Center for Computational Toxicology and
Exposure

Journal of Cheminformatics

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Methodology | [Open Access](#) | Published: 30 August 2018

"MS-Ready" structures for non-targeted high-resolution mass spectrometry screening studies

[Andrew D. McEachran](#) , [Kamel Mansouri](#), [Chris Grulke](#), [Emma L. Schymanski](#), [Christoph Ruttkies](#) & [Antony J. Williams](#) 

Journal of Cheminformatics 10, Article number: 45 (2018) | [Cite this article](#)

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Our Curation Process

Computational Toxicology 12 (2019) 100096



Contents lists available at [ScienceDirect](#)

Computational Toxicology

journal homepage: www.elsevier.com/locate/comtox



EPA's DSSTox database: History of development of a curated chemistry resource supporting computational toxicology research



Christopher M. Grulke^a, Antony J. Williams^a, Inthirany Thillanadarajah^b, Ann M. Richard^{a,*}

Journal of Cheminformatics

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Research article | [Open Access](#) | Published: 08 March 2018

OPERA models for predicting physicochemical properties and environmental fate endpoints

[Kamel Mansouri](#) , [Chris M. Grulke](#), [Richard S. Judson](#) & [Antony J. Williams](#)

Journal of Cheminformatics **10**, Article number: 10 (2018) | [Cite this article](#)

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TEST – Toxicity Estimation Software Tool

<https://www.epa.gov/chemical-research/toxicity-estimation-software-tool-test>

Toxicity Estimation Software Tool (TEST)

On this page:

- [QSAR Methodologies](#)
- [What's New in Version 4.2.1?](#)
- [Prior Version History](#)
- [System Requirements](#)
- [Installation Instructions](#)
- [Publications](#)
- [Get Email Alerts](#)

The Toxicity Estimation Software Tool (TEST) was developed to allow users to easily estimate the toxicity of chemicals using Quantitative Structure Activity Relationships (QSARs) methodologies. QSARs are mathematical models used to predict measures of toxicity from the physical characteristics of the structure of chemicals (known as molecular descriptors). Simple QSAR models calculate the toxicity of chemicals using a simple linear function of molecular descriptors:

Ask a Technical Expert

Got a question about our research model?
Want to give us feedback? Contact a technical expert about [TEST](#).



Structure/Substructure Search

atrazine

Select properties to predict

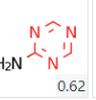
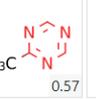
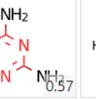
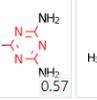
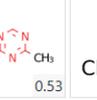
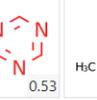
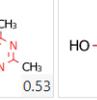
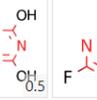
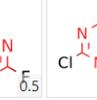
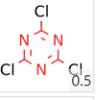
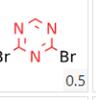
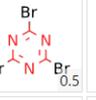
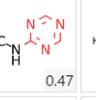
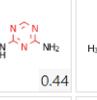
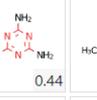
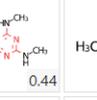
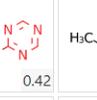
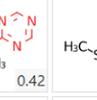
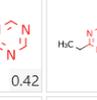
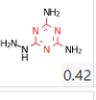
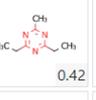
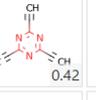
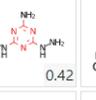
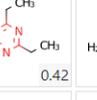
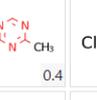
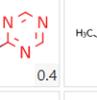
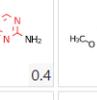
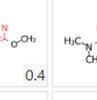
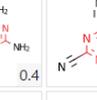
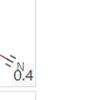
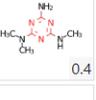
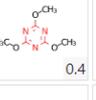
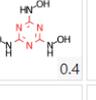
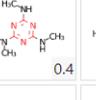
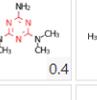
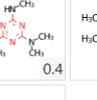
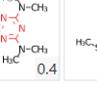
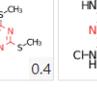
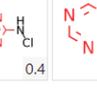
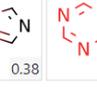
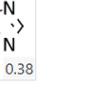
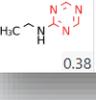
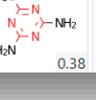
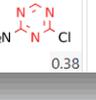
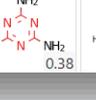
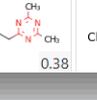
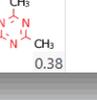
T.E.S.T. **18** OPERA

- Exact
- Substructure
- Similarity
- Molecular Formula
- Molecular Weight

Filter by elements (enter comma separated list e.g. C,F,H) include

Filter by elements (enter comma separated list e.g. C,F,H) exclude

Search result **2540** Show Isotopically Labeled Charged Salts or Mixtures Sort Similarity

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 0.4	 0.4	 0.4	 0.4	 0.4	 0.4	 0.4	 0.4	 0.4	 0.4	 0.38
 0.38	 0.38	 0.38	 0.38	 0.38	 0.38	 0.38	 0.38	 0.38	 0.38	 0.38



Spectral Searching against Predicted Fragmentation

Non Target Analysis Prototype

Mass Search

 Da

Molecular Formula Search

Mass or Formula must be entered before searching spectrum

Ionization Type

ESI+
ESI+
ESI-
EI

Spectra Input

304.1332052	11.6199475
198.0913404	7.308439899
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196.0756904	5.289463115
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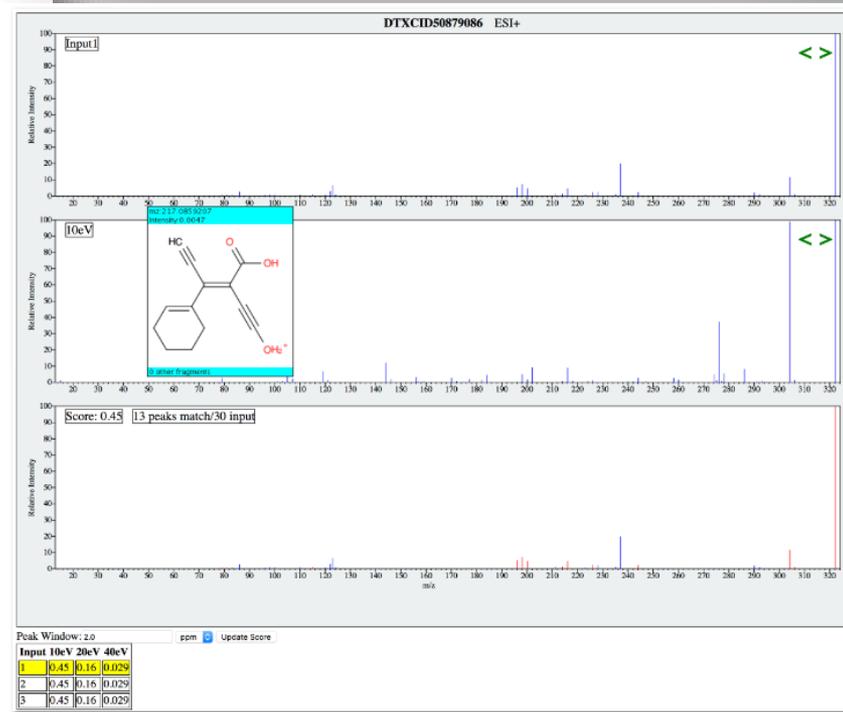
Peak Match Window:

Data Descriptor | [OPEN](#) | Published: 02 August 2019

Linking *in silico* MS/MS spectra with chemistry data to improve identification of unknowns

Andrew D. McEachran , Ilya Balabin, Tommy Cathey, Thomas R. Transue, Hussein Al-Ghoul, Chris Grulke, Jon R. Sobus & Antony J. Williams

Scientific Data **6**, Article number: 141 (2019) | [Download Citation](#) ↓





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