

Consensus ranking and fragmentation prediction for identification of unknowns in high resolution mass spectrometry

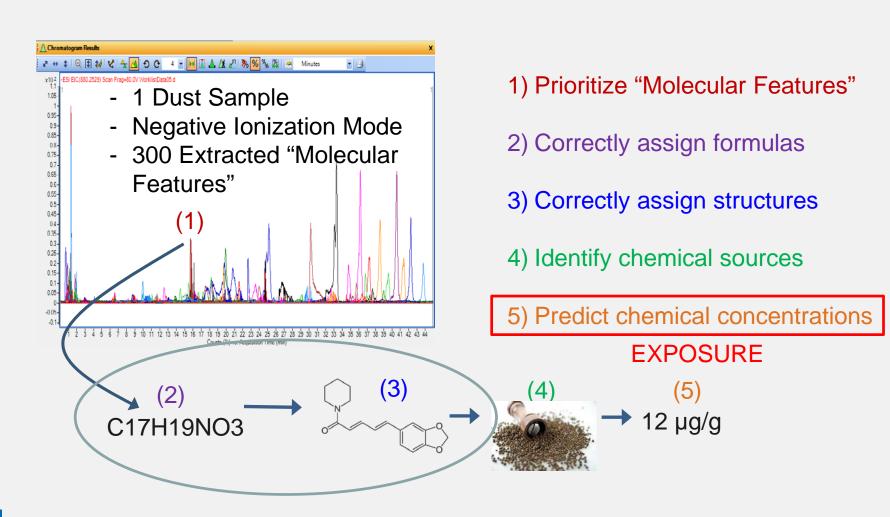
Andrew D. McEachran Hussein Al-Ghoul, Ilya Balabin, Tommy Cathey, Alex Chao, Jon Sobus, and Antony J. Williams

AGRO 107

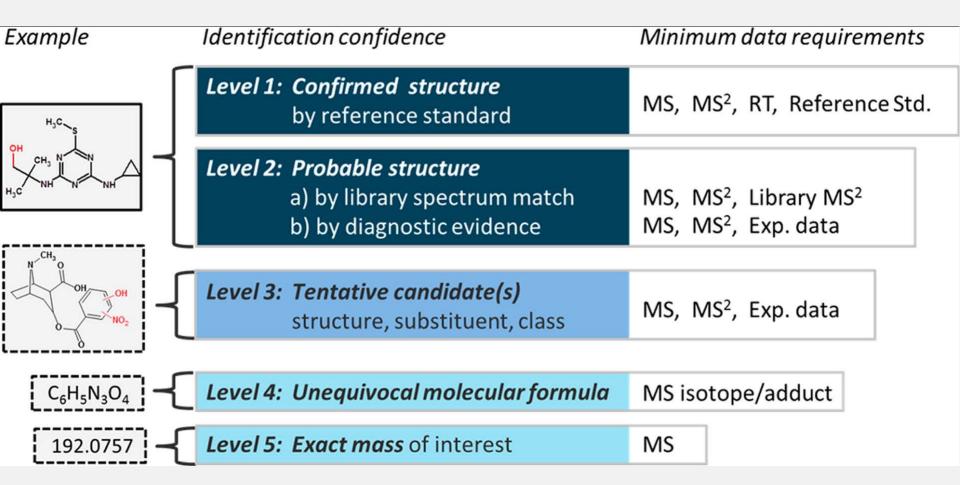
The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA



General Goals of SSA/NTA

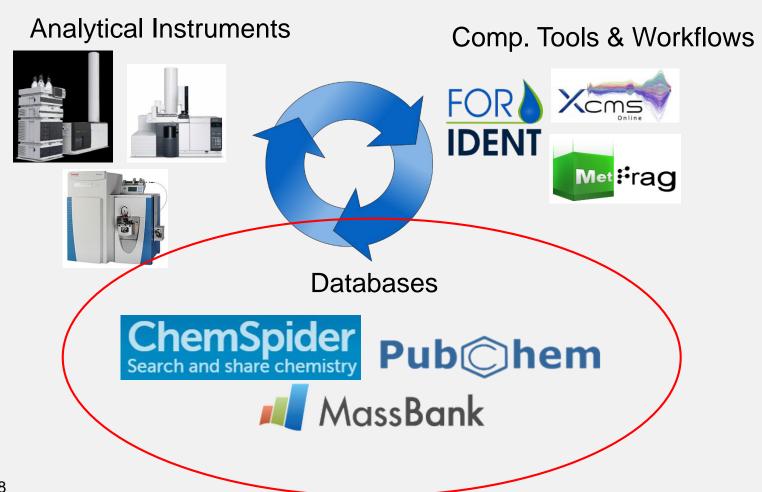






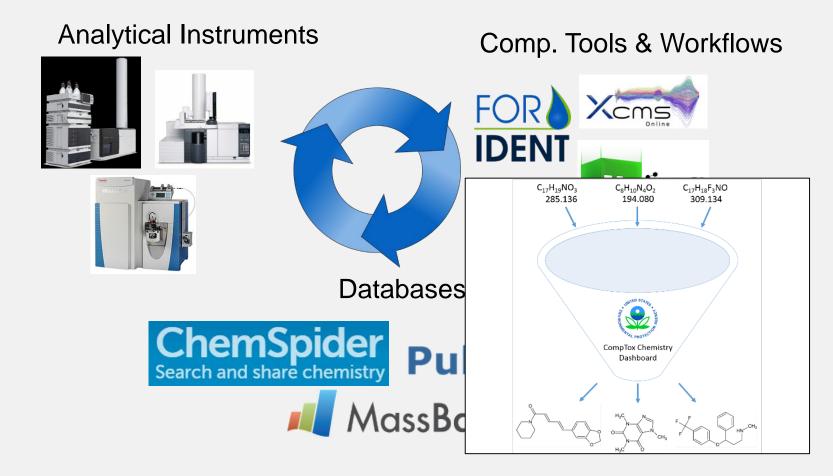


The General Approach



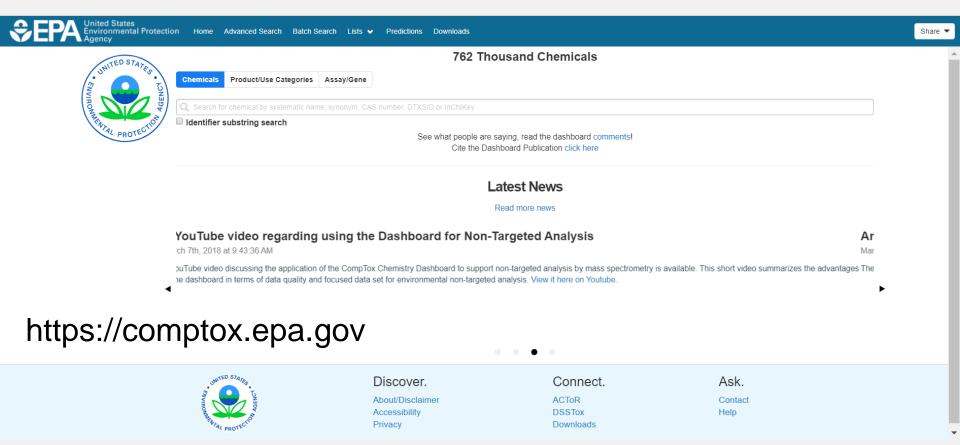


The General Approach





CompTox Dashboard

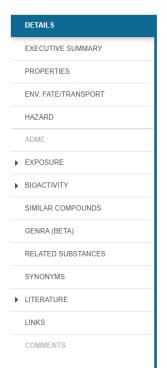


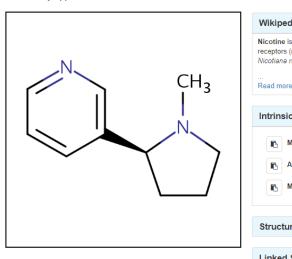


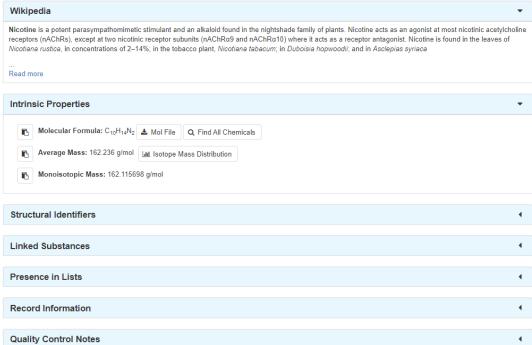
Nicotine

54-11-5 | DTXSID1020930

Searched by Approved Name.













Home Advanced Search Batch Search Lists ♥ Predictions Downloads

Property

Summary

▲ Download ▼

Columns V



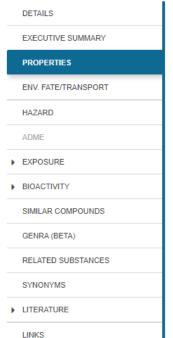
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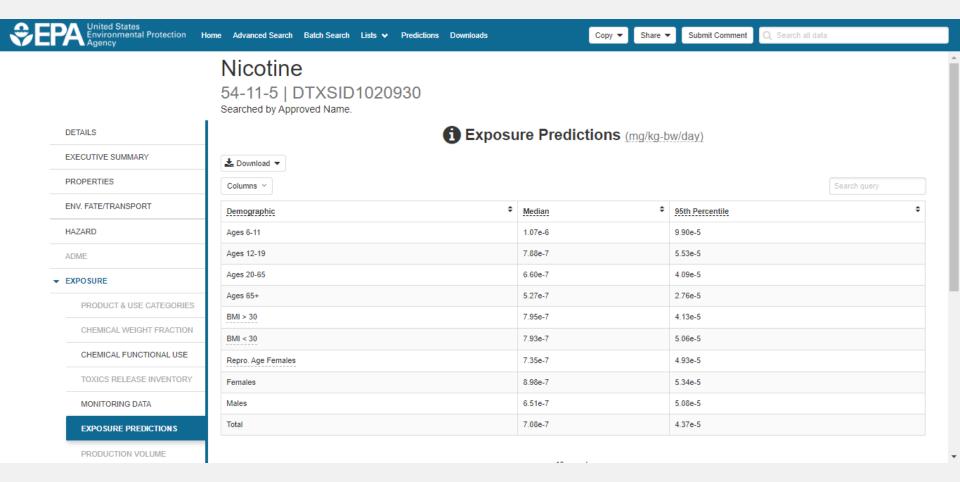
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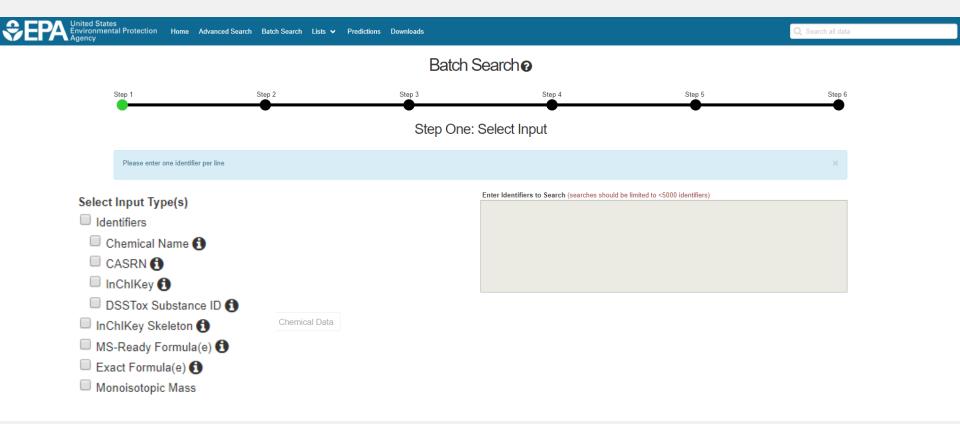
Summary

Property	Experimental average \$	Predicted average \$	Experimental median	Predicted median *	Experimental range \$	Predicted range \$	<u>Unit</u>
LogP: Octanol-Water	1.17 (1)	0.751		0.821	1.17	3.85e-2 to 1.18	
Melting Point	-79.0 (3)	12.4	-79.0	13.4	-79.0	-34.4 to 57.3	°C
Boiling Point	247 (2)	249	247	248	247	244 to 254	°C
Vapor Pressure	3.80e-2 (1)	1.70e-2		1.76e-2	3.80e-2	2.39e-3 to 3.03e-2	mmHg
Water Solubility	6.16 (1)	3.74		4.51	6.16	8.00e-2 to 6.63	mol/L
Flash Point	-	99.8		99.8	-	97.9 to 102	°C
Surface Tension	-	38.6		38.6	-	37.7 to 39.6	dyn/cm
Index of Refraction	-	1.54			-	1.54	
Molar Refractivity	-	49.3			-	49.3	cm^3
Polarizability	-	19.5			-	19.5	Å^3



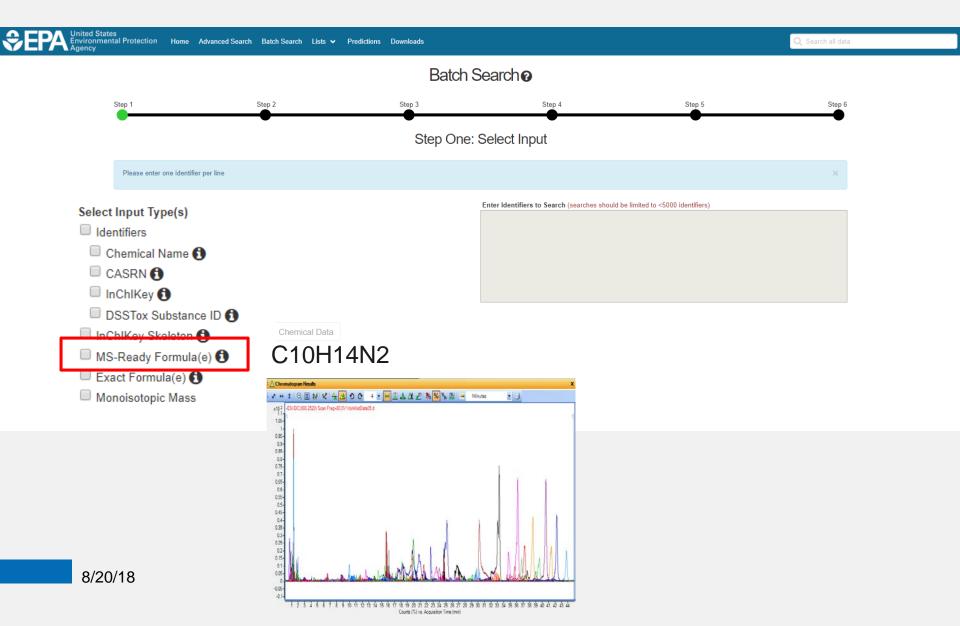






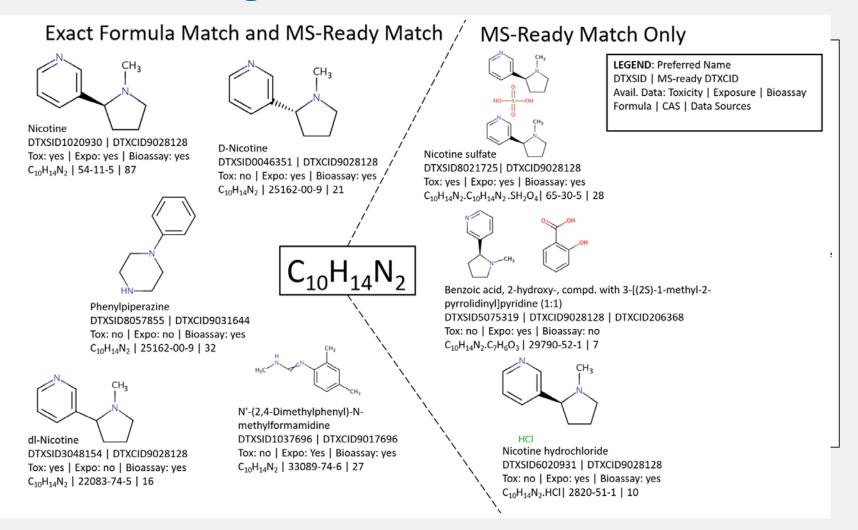


Batch Search for SSA/NTA





MS-Ready Structures improve database searching





Data Source Ranking of "known unknowns"

- Mass and/or formula unknown to a researcher, contained within a reference database
- Most likely candidate chemicals have the most references/sources

C14H22N2O3 266.16304



Chemical Reference Database



Sorted candidate structures



Initial Data Source Ranking in ChemSpider

AS MS

C American Society for Mass Spectrometry, 2011

J. Am. Soc. Mass Spectrom. (2012) 23:179–185DOI: 10.1007/s13361-011-0265-y

Adopted by NTA researchers around the world

RESEARCH ARTICLE

Identification of "Known Unknowns" Utilizing Accurate Mass Data and ChemSpider

Table 1. Searching ChemSpider by Elemental Composition then Sorting by Number of Associated References

Class of compounds	Number compounds in class	Position of compound sorted in descending order by number of references					
		#1	#2	#3	#4	#5	>#5
Drugs	45	43	1	1			
Pesticides	8	7	1				
Toxins	2	2					
Polymer antioxidants	15	15					
Polymer UV stabilizers	10	8	1	1			
Polymer clarifying agent (Irgaclear DM)	1						1(14)
Polyurethane additives	4	2	1			1	
Natural products	3	2		1			
Herbicide (clofibric acid)	1	1					
Artificial sweetener (sucralose)	1	1					
Total compounds ChemSpider	90	81	4	3		1	1
Total compounds CAS Registry [1]	90	84	4	1		1	





RAPID COMMUNICATION

Identifying known unknowns using the US EPA's CompTox Chemistry Dashboard

Andrew D. McEachran 1 · Jon R. Sobus 2 · Antony J. Williams 3

 On same 162 chemicals, Dashboard outperforms ChemSpider

	Mass-based searching		Formula-based searching		
	Dashboard	ChemSpider	Dashboard	ChemSpider	
Average rank position Percent in #1 position	1.3 85%	2.2 ^a 70%	1.2 88%	1.4 80%	

^a Average rank in ChemSpider shown here does not include an outlier where the rank was 201, when added the average rank position is 3.5



Additional Data Streams to Improve Identifications

- US EPA CompTox Dashboard Data Sources (DS)
- PubChem Data Source Count
- PubMed Reference Count

Pub©hem

- Presence in STOFF-IDENT Database
- Predicted Environmental Media Occurrence
- OPERA PhysChem Properties
- NORMAN Network Priority List

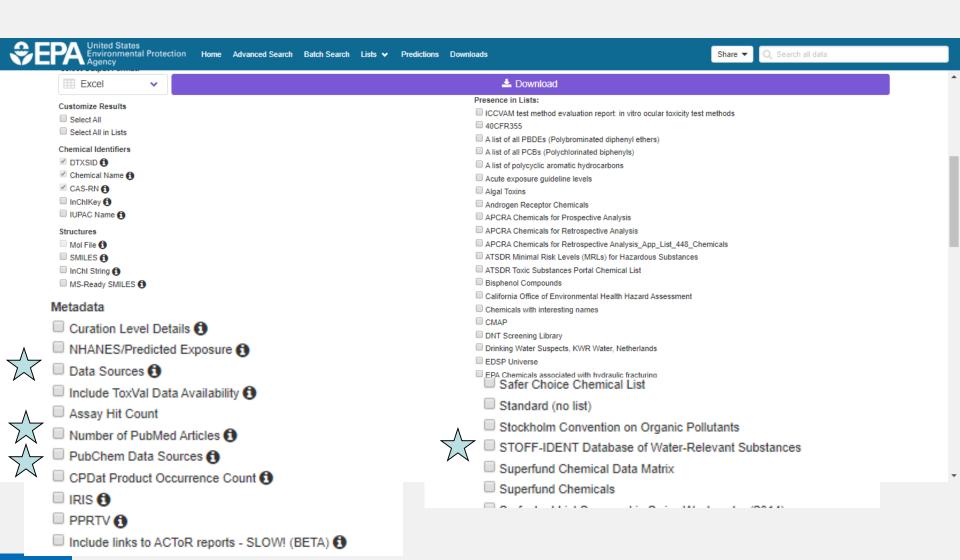








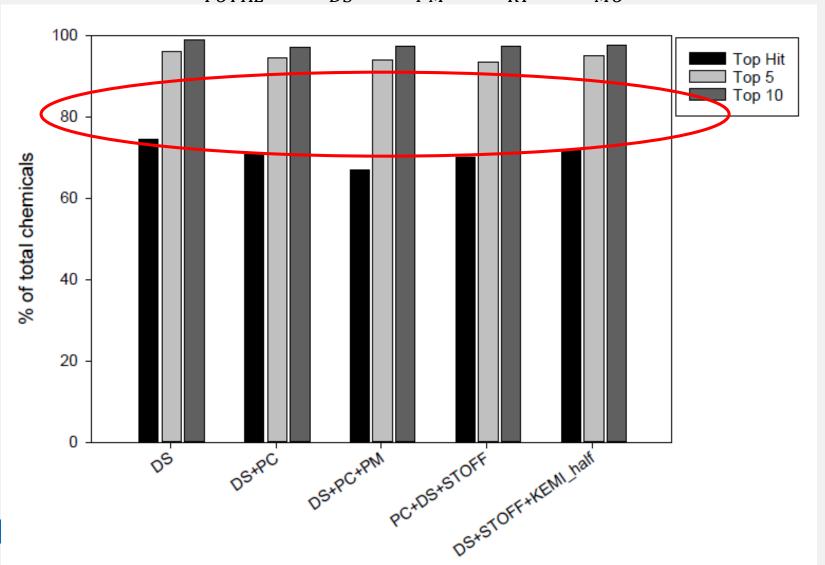
All available via Batch Search:





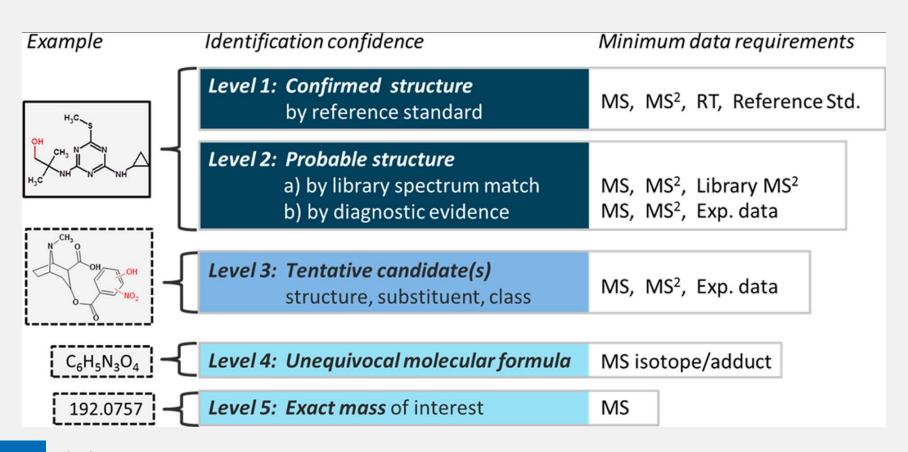
Identification ranks for 1783 chemicals using multiple data streams

$$SC_{TOTAL} = SC_{DS} + SC_{PM} + SC_{RT} + SC_{MO} + \cdots$$





Metadata is critical, but need structural confirmation to increase confidence

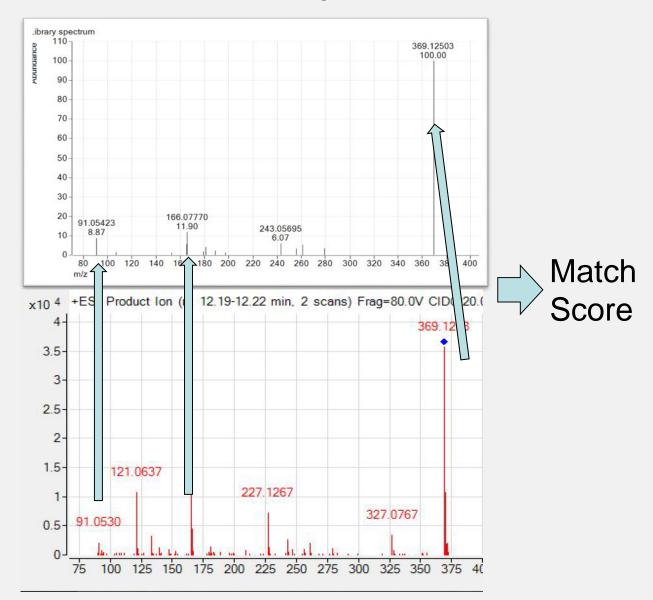




MS/MS Spectral Matching for Identification

Library Fragmentation Spectra (20eV)

Observed Fragmentation Spectra (20eV)







Metabolomics

February 2015, Volume 11, Issue 1, pp 98-110 | Cite as

Competitive fragmentation modeling of ESI-MS/MS spectra for putative metabolite identification

Authors and affiliations

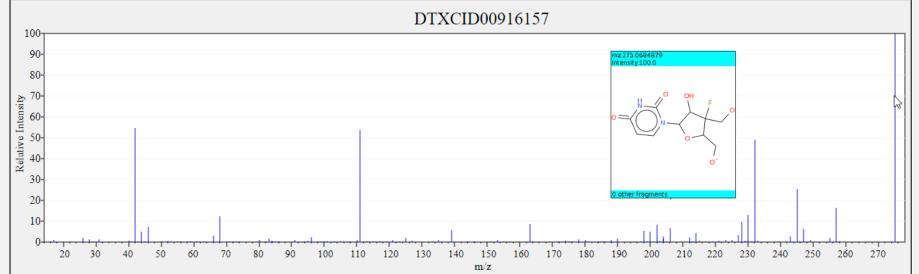
Felicity Allen , Russ Greiner, David Wishart

CFM-ID

- Fragmentation prediction for identification in HRMS
- Open source code allows for MS/MS spectra prediction for ESI+, ESI-, and EI
- Predictions generated and stored for >700,000 structures, to be accessible via CompTox Dashboard
- Python code to pull matches and score experimental vs predicted spectra
- Cosine dot product match score calculation



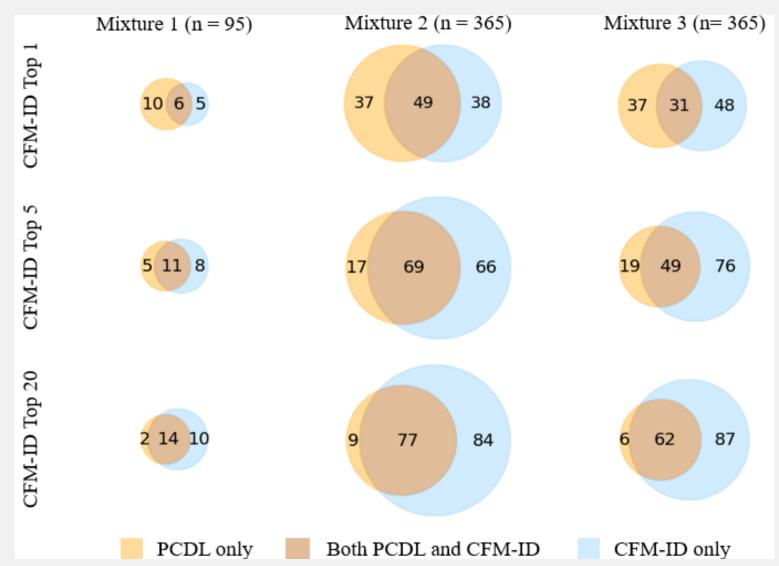




```
97 44.04947561 CC=[NH2+]
98 112.0393049 C=C(C([CH2+])=N)C(=0)0
# Date/time: 02/13/18 22:23:52
# CFM-ID version: 2.0 snapshot 10/23/2017
# DTXCID: DTXCID40539667
# SMILES: CCOC(=0)C1=C(C)N=C2C=C(C)C(C)=CC2=C1
# MASS: 243.125928791
# FORMULA: C15H17NO2
# INCHI KEY: PKWJEOYUFQXCBY-UHFFFAOYSA-N
energy0
15.02292652 0.2213297725 4 (0.22133)
27.02292652 0.1671394176 13 (0.16714)
29.00219107 0.003415204535 26 (0.0034152)
29.03857658 0.7931575846 14 (0.79316)
41.00219107 0.05449899314 52 (0.054499)
42.03382555 0.002436421336 104 (0.0024364)
43.01784114 0.03020513753 53 (0.030205)
44.99710569 0.002435326653 84 (0.0024353)
45.0334912 0.045173947 54 (0.045174)
46.06512568 0.00275766249 105 (0.0027577)
47.01275576 0.001588587881 83 (0.0015886)
47.04914126 0.7597376869 55 (0.75974)
51.02292652 0.0008416543192 41 (0.00084165)
57.0334912 0.002233145503 57 (0.0022331)
65.00219107 0.0008022807615 30 (0.00080228)
65.03857658 0.005921490873 43 (0.0059215)
67.05422664 0.01637516089 112 (0.016375)
68.99710569 0.0308626766 80 (0.030863)
69.06987671 0.07637646561 127 (0.076376)
71.01275576 0.01301877006 81 (0.013019)
73.02840582 0.09231084238 82 (0.092311)
75.04405588 0.03569749353 85 (0.035697)
103.0542266 0.05442462493 70 (0.054425)
105.0698767 0.07415736715 96 (0.074157)
117.0698767 0.04642433142 94 (0.046424)
```



Predicted MS/MS spectra provide greater coverage than empirical libraries







Evaluating on CASMI 2016

- Critical Assessment of Small Molecule Identification
 - -Training data= 312 peak lists (from 285 substances)
 - 234 MS/MS in positive mode
 - 58 in negative mode
 - -Challenge Data= 208 peak lists (from 188 substances)
 - 127 in positive mode
 - 81 in negative mode
- Precursor ion search window= 15 ppm
- Fragment ion match threshold= 0.02 Da
- Candidates limited to Dashboard results within precursor ion search window





CASMI 2016 Contest Challenge Set (n=208)

CFM-ID only

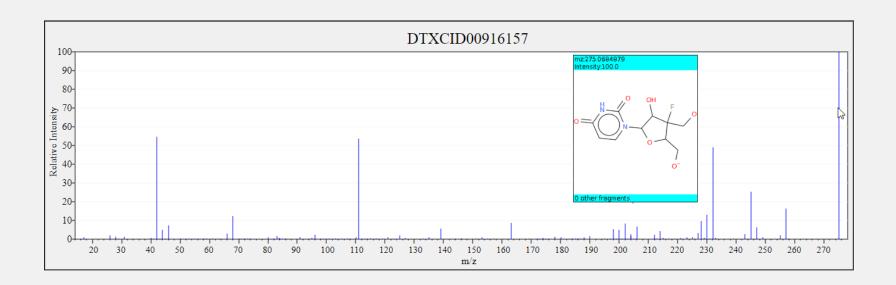
CFM-ID +DSSTox Data Sources

	# Identified	% of Total
#1 Hits	89	43%
Top 5	154	74%
Top 10	174	84%
Top 20	190	91%

	# Identified	% of Total
#1 Hits	154	74%
Top 5	195	94%
Top 10	198	95%
Top 20	202	97%



Access via CompTox Dashboard



Data available for download after publication



8/20/1

Search

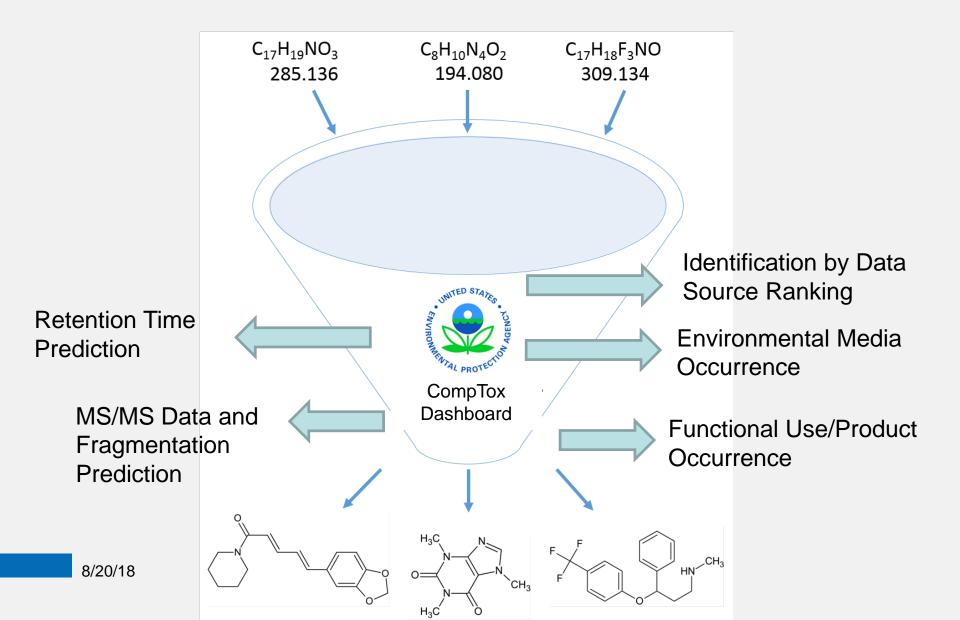
Mockup, work in progress....

Non Target Analysis Prototype

	arch				
±	Min/Max				
322.14431847	76	Da	±	5	Da ppm
Molecular	r Formula	Search			
Molecular For	mula				
Ionization					
	•				
Single	Energy	Multiple			
304.1332052 198.0913404 123.0440559 196.0756904	11.6199475 7.306439699 6.538348292 5.269463115 4.700461978	Multiple			



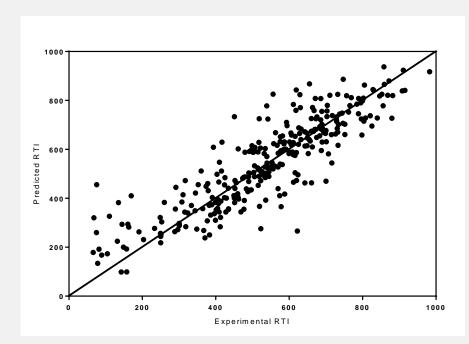
Dashboard in NTA Workflows





Future Directions

- Combined data visualization
- Retention time index (RTI) predictions
- Ongoing expansion of the database
- Integration to public MS databases





Conclusions

- Databases are effective resources in SSA/NTA
- CompTox Dashboard provides access to chemistry data for >760,000 chemical substances
- Predicted MS/MS spectra linked within the CompTox Dashboard further enhances effectiveness and increases confidence in identifications



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Seth Newton

External Collaborators

Emma Schymanski- Univ.

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Christoph Ruttkies- IPB,

Halle

Kamel Mansouri- ILS, Inc

*ORISE Research Participant



Questions?

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- http://orcid.org/0000-0003-1423-330X
- Associated presentations:
 - AGRO 29: Leveraging chemistry data to improve exposure analyses using the EPA's CompTox Chemistry Dashboard
 - ANYL 100: Developing tools for high resolution mass spectrometry-based screening via the EPA's CompTox Chemistry Dashboard
 - ENVR 152: EPA Comptox Chemistry Dashboard as a data integration hub for environmental chemistry data