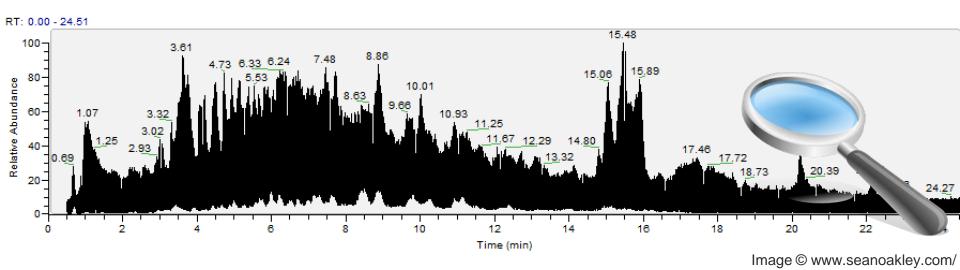


Finding Small Molecules in Big Data





Emma Schymanski
Luxembourg Centre for Systems Biomedicine (LCSB),
University of Luxembourg
Email: emma.schymanski@uni.lu

Antony J. Williams

National Centre for Computational Toxicity (NCCT), US Environmental Protection Agency (US EPA), NC, USA





Small molecules ... big problems?

Status quo of small molecules:

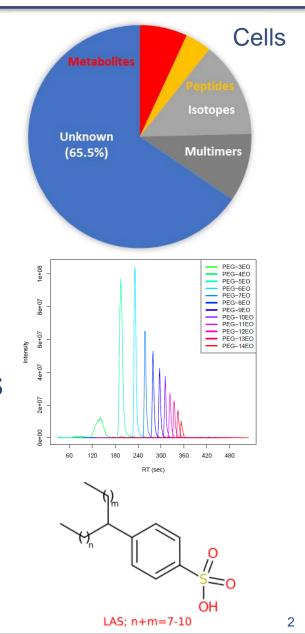
- How many are in compound databases?
- How many could there be?
- How many are in spectral libraries?
- Mind the Gap!

Exchanging "expert knowledge"

- Suspect Lists in Europe
- Live, retrospective screening & untargeted MS

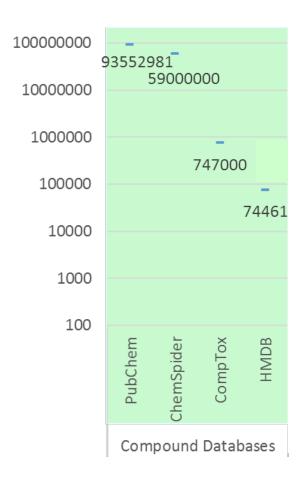
Tackling Complex Structures

- Exchanging Information on Unknowns
- …and how Open Science helps!



Searching for Small Molecules ...

Compound Databases



PubChem: >95 million https://pubchem.ncbi.nlm.nih.gov/

ChemSpider: >60 million http://www.chemspider.com/

CompTox Chemicals Dashboard: >761 000 https://comptox.epa.gov/dashboard/

Human Metabolome DB (HMDB): >115 000 http://www.hmdb.ca/

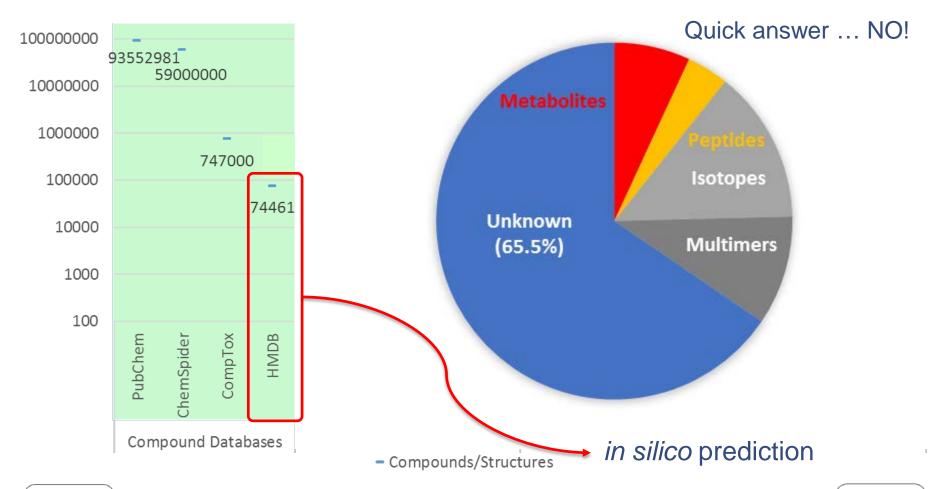
- Compounds/Structures





Searching for Small Molecules ...

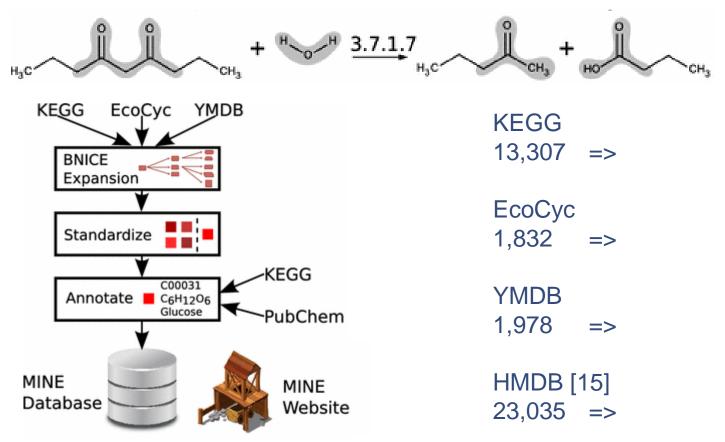
Compound Databases ... isn't 95 million enough?





Searching for More Small Molecules ...

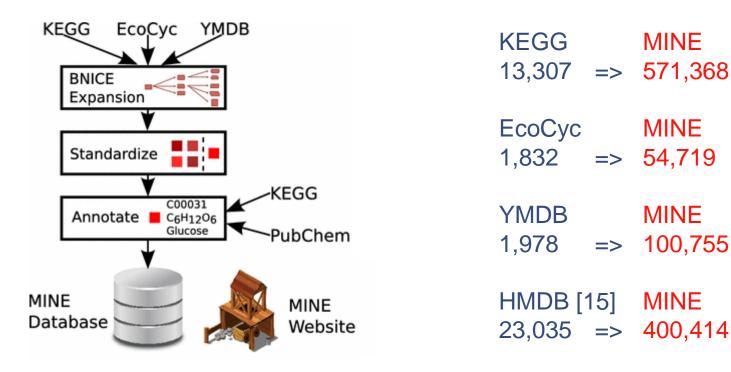
In silico metabolite prediction – example of MINE (2015)





Searching for More Small Molecules ...

- *In silico* metabolite prediction example of MINE (2015)
 - First generation only ... combinatorial explosion!



Speculation ...



MINE

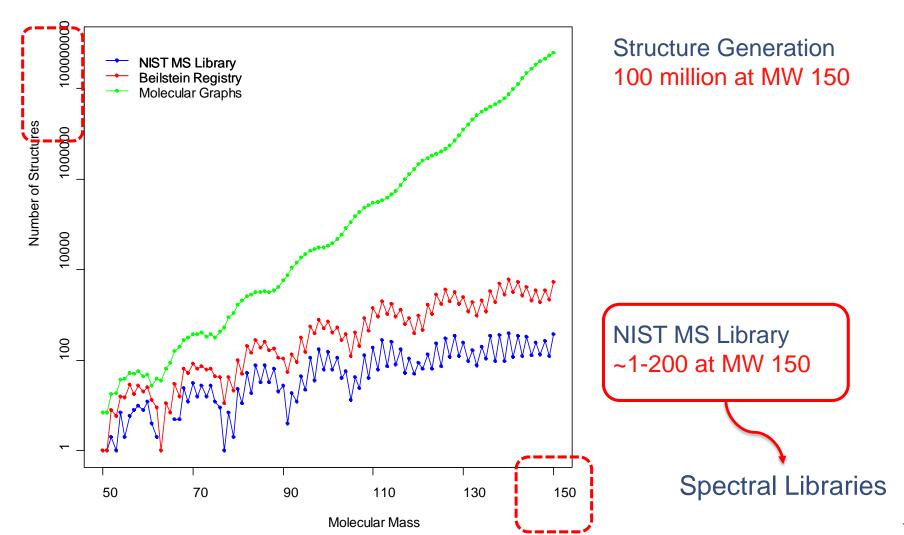
=> 1.6 billion ... first generation only?!?!



Searching for EVEN MORE Small Molecules ...

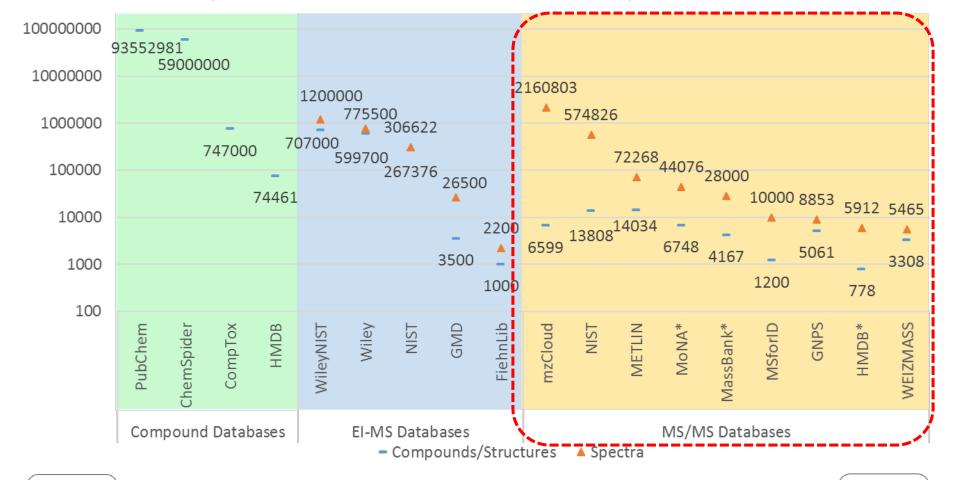
Structure Generation

But of course most of these do not exist



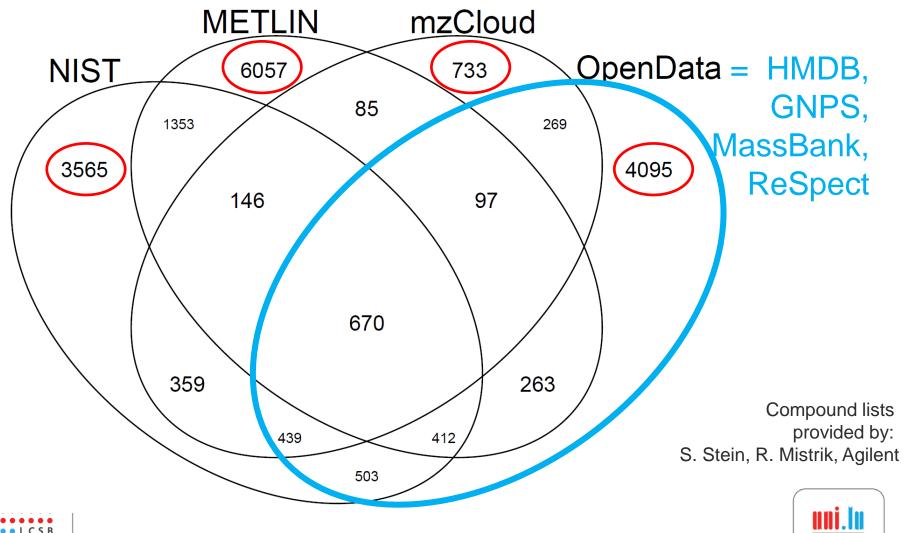
Searching for Small Molecules in Spectral Libraries

- ... to find what is "on record"...
 - Too many different MS/MS libraries (and they are still too small)



Do we need all these libraries?

Yes ... most libraries still have many unique entries

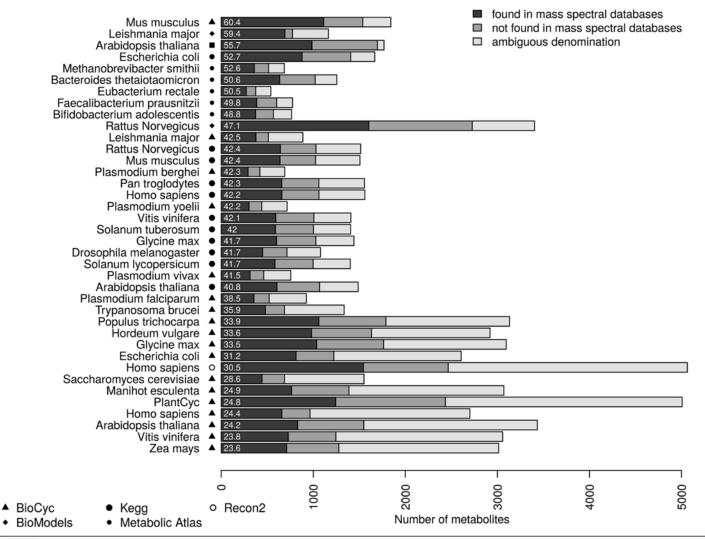


Mind the Gap!

AraGEM

BIGG

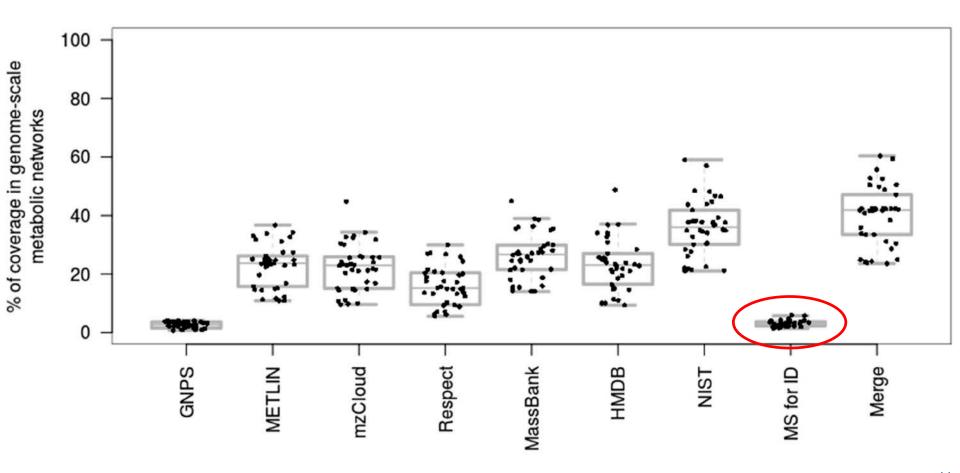
 Only 23-60 % of (defined) metabolites in Genome-Scale Metabolic Networks are covered by (combined!) Mass Spectral Libraries



10

Mind the Gap!

- Best library to choose depends highly on your dataset
 - Example: MSforID (https://msforid.com/) is poor for metabolic networks but great for forensic toxicology!



SPectraL hASH (SPLASH) – Search between libraries

http://splash.fiehnlab.ucdavis.edu/

splash10 - 0002 - 0900000000 - b112e4e059e1ecf98c5f

[version] - [top10] - [histogram] - [hash of full spectrum]



splash10-0002-0900000000-b112e4e059e1ecf98c5f



Human Metabolome Database: LC-MS/MS Spectrum - LC-ESI-QTOF ... www.hmdb.ca/spectra/ms_ms/5464 ▼

... Spectrum - LC-ESI-QTOF (UPLC Q-Tof Premier, Waters) 30V, Positive. Splash Key: splash10-0002-0900000000-b112e4e059e1ecf98c5f View in MoNA ...

Human Metabolome Database: Showing metabocard for Caffeine ... www.hmdb.ca/metabolites/HMDB01847 ▼



splash10-0uxr-0973000000-87d07ddd2ed24b9598d7



DrugBank: Codeine

www.drugbank.ca/drugs/DB00318 ▼

... 60V, Positive, **splash10-0uxr-0973000000-87d07ddd2ed24b9598d7**, View in MoNA. MS, Mass Spectrum (Electron Ionization), splash10-01ot-3950000000- ...

Codeine Mass Spectrum - MassBank

massbank.eu/MassBank/jsp/Dispatcher.jsp?type=disp&id=EA278005&site=31 ▼ PK\$SPLASH: splash10-0uxr-0973000000-87d07ddd2ed24b9598d7 PK\$ANNOTATION: m/z tentative_formula_count mass error(ppm) 58.0651 ...

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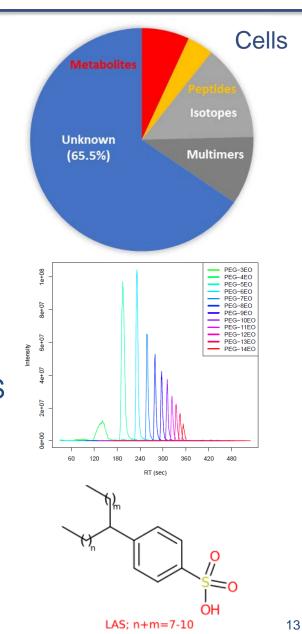
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Exchanging "expert knowledge"

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- Live, retrospective screening & untargeted MS

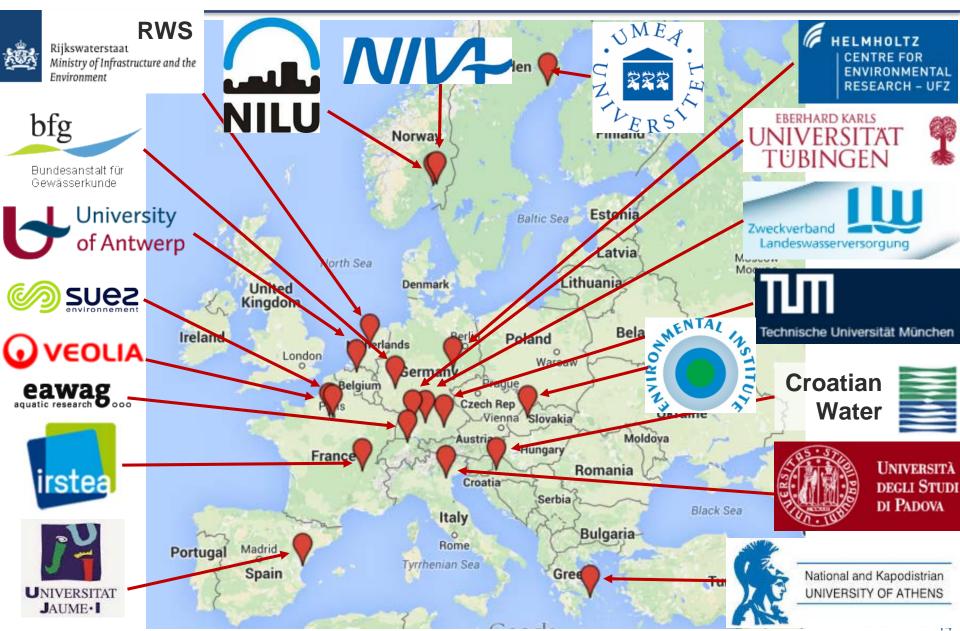
Tackling Complex Structures

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- ...and how Open Science helps!



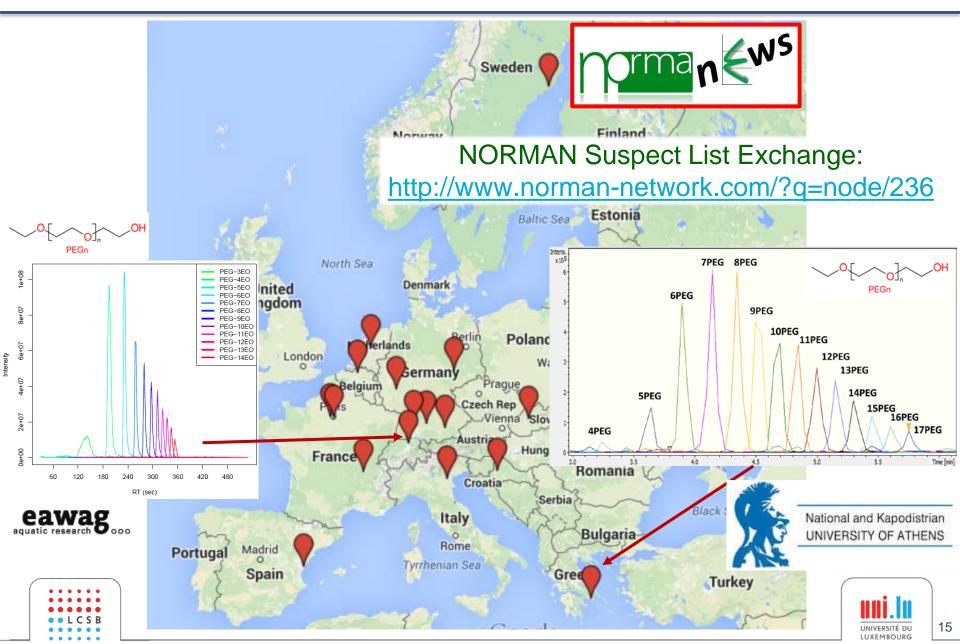
2015: European Non-target Screening Trial





European (World-)Wide Exchange of Suspects

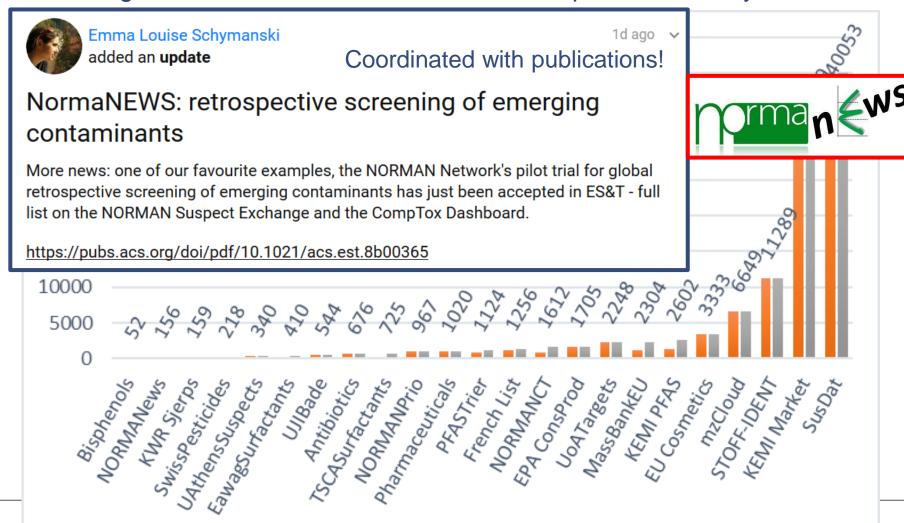




NORMAN Suspect Exchange Lists



- http://www.norman-network.com/?q=node/236
- 21 lists available ... specialist collections to market lists
 - Integrated in NORMAN Databases & CompTox Chemistry Dashboard





man ews on the CompTox Chemicals Dashboard EPA

https://comptox.epa.gov/dashboard/chemical_lists/normanews

Normanews: Norman Early Warning System

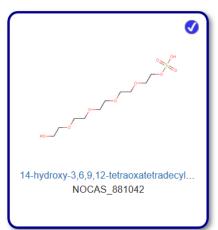
Search NORMANEWS Chemicals

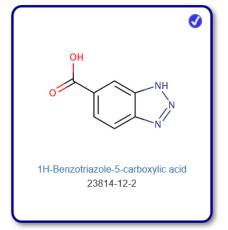
List Details

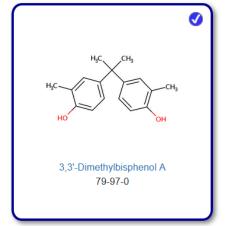
Description: The Norman Early Warning System (NormanEWS) is a pilot network designed to investigate the spatial and temporal distribution of newly identified contaminants of emerging concern in environmental samples through performing retrospective suspect screening on HRMS data acquired using different instrumental platforms and data processing software. The NormaNEWS pilot study was performed through recruiting eight reference laboratories with available archived HRMS data with the goal of exploring the potential of an early warning network to rapidly establish the occurrence of newly-identified contaminants of emerging concern across Europe and beyond, through the use of retrospective suspect screening employing HRMS. The pilot study was referred to as the Norman Early Warning System, abbreviated to NormanEWS

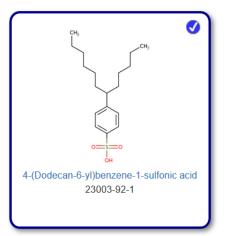
Number of Chemicals: 131

4 of 131 chemicals selected Download / Send Unselected Sort by: Name Deselect all









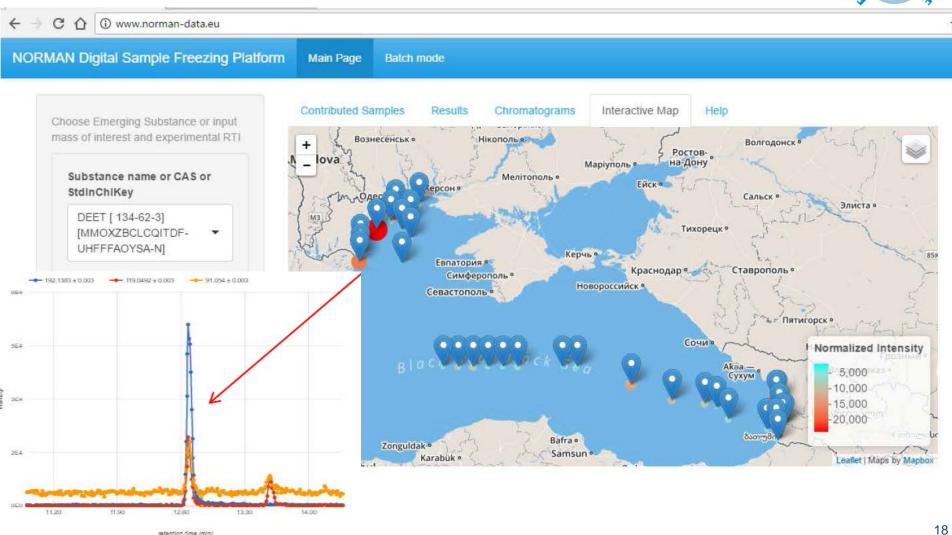
Q

NORMAN Digital Sample Freezing Platform



"Live" retrospective screening of known and unknown chemicals in European samples (various matrices)





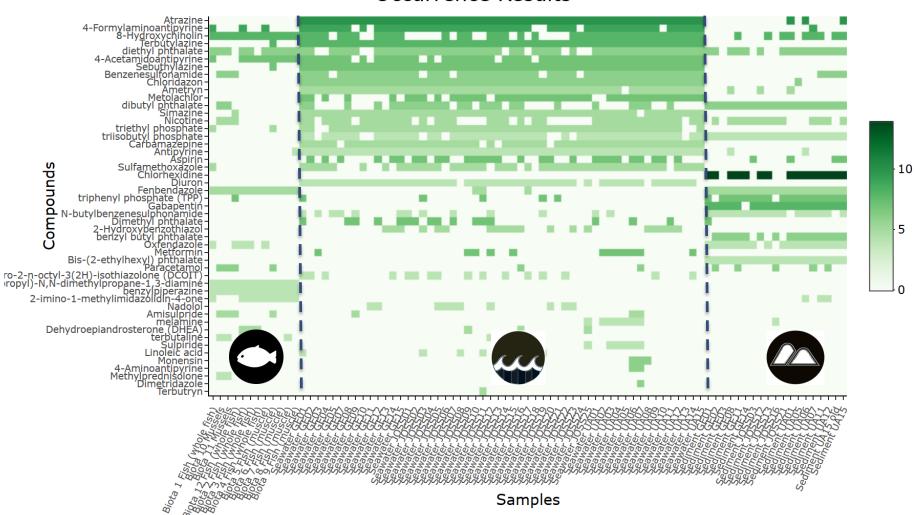
NORMAN Digital Sample Freezing Platform



Retrospective screening of REACH chemicals in Black Sea samples (various matrices)

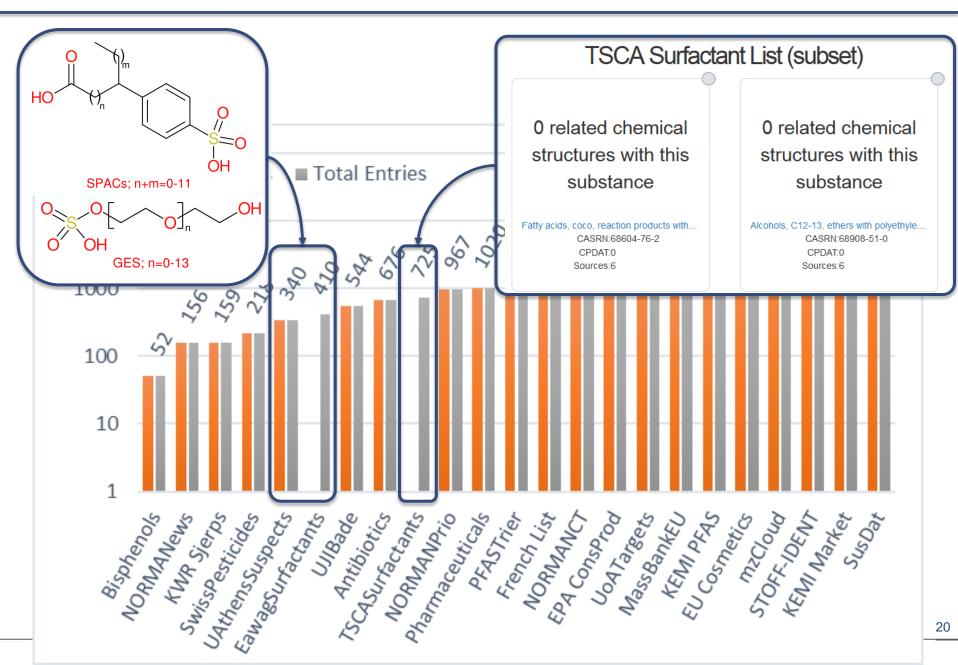


Occurrence Results



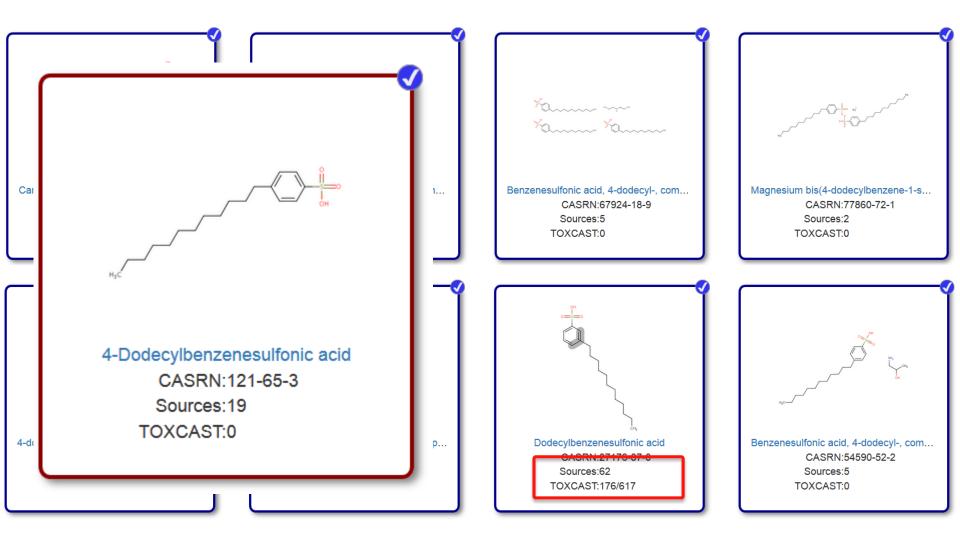
Challenge 1: Not all suspects have structures!





Challenge 2: Mass Spec "sees" ONE part at a time

O What do all these chemicals have in common?



Small molecules ... big problems?

Status quo of small molecules:

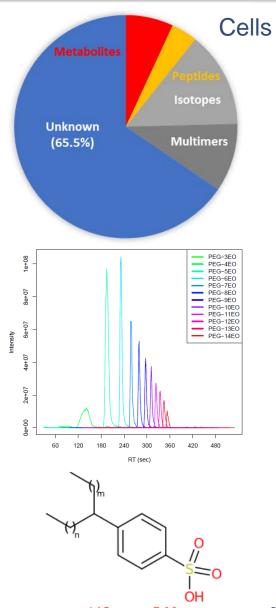
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Exchanging "expert knowledge"

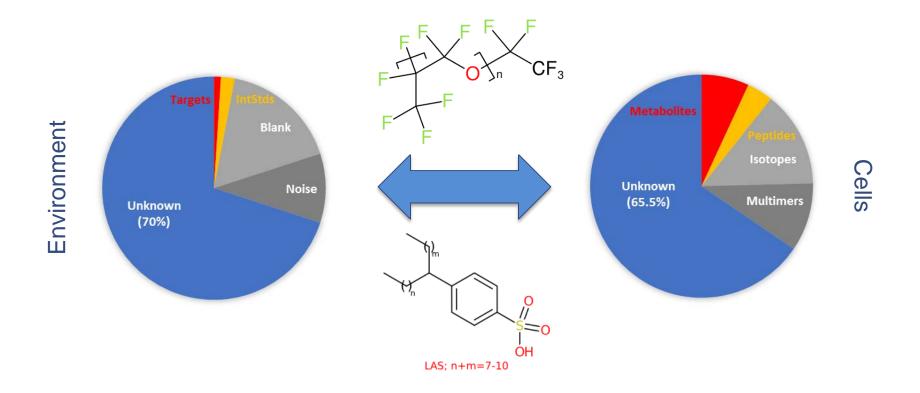
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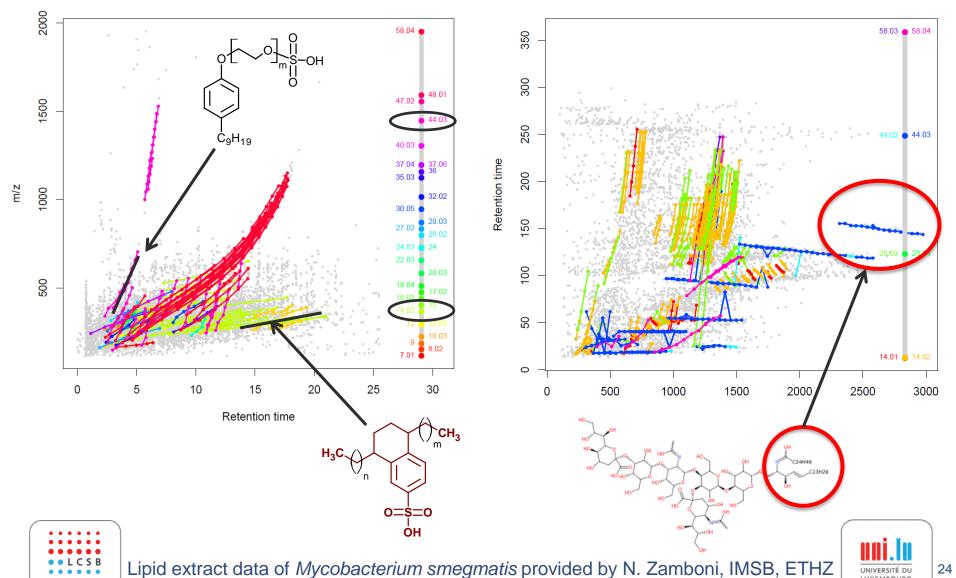
We still have many unknowns ...



...and many are interconnected by mass

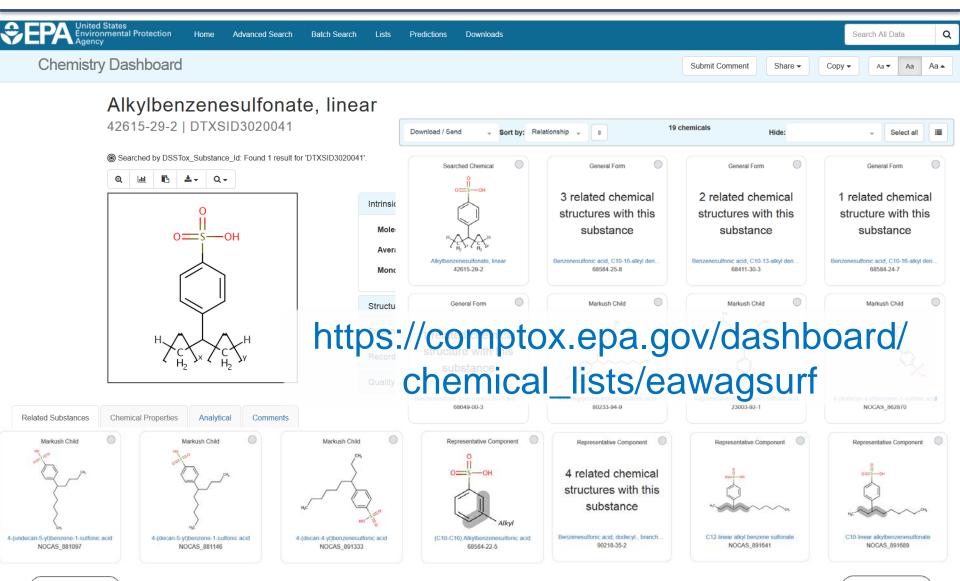


Homologous Series in environmental and biological samples



New Ways to Store and Access Homologues



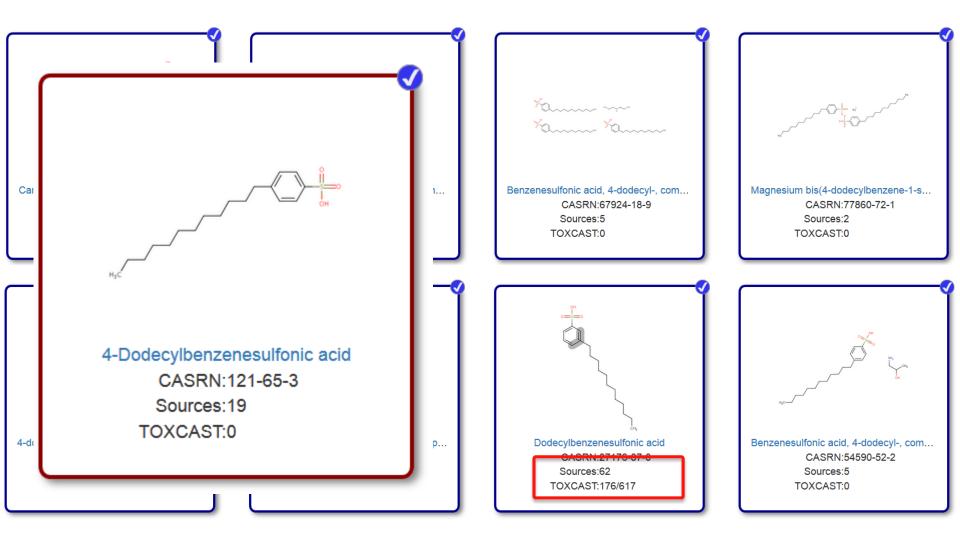


https://comptox.epa.gov/dashboard/dsstoxdb/results?search=DTXSID3020041 https://comptox.epa.gov/dashboard/chemical_lists/eawagsurf

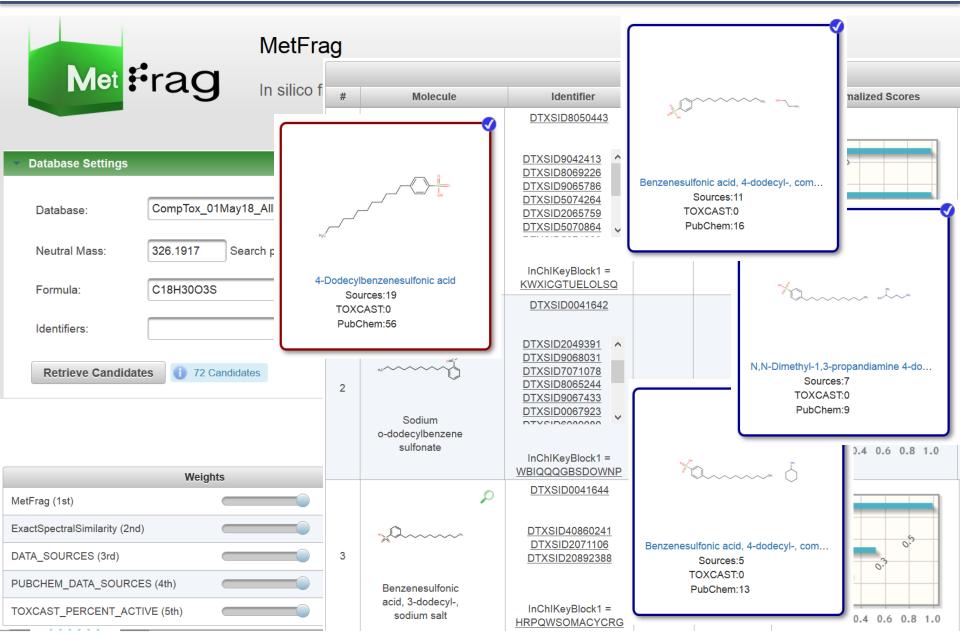


RECAP: Mass Spec "sees" ONE part at a time

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MS-Ready: Accessing Data from Salts and Mixtures



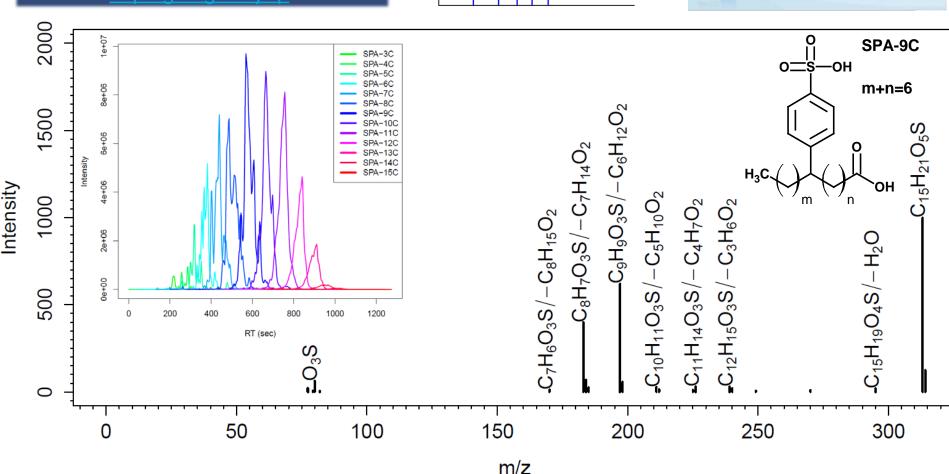
McEachran et al. 2018 J. Cheminformatics 10:45 DOI: 10.1186/s13321-018-0299-2 and https://msbi.ipb-halle.de/MetFragBeta/

Annotated Spectra of Homologues in MassBank.EU



RMass Bank





www.massbank.eu

https://github.com/MassBank/RMassBank/

ACCESSIONS (LAS, SPACs):

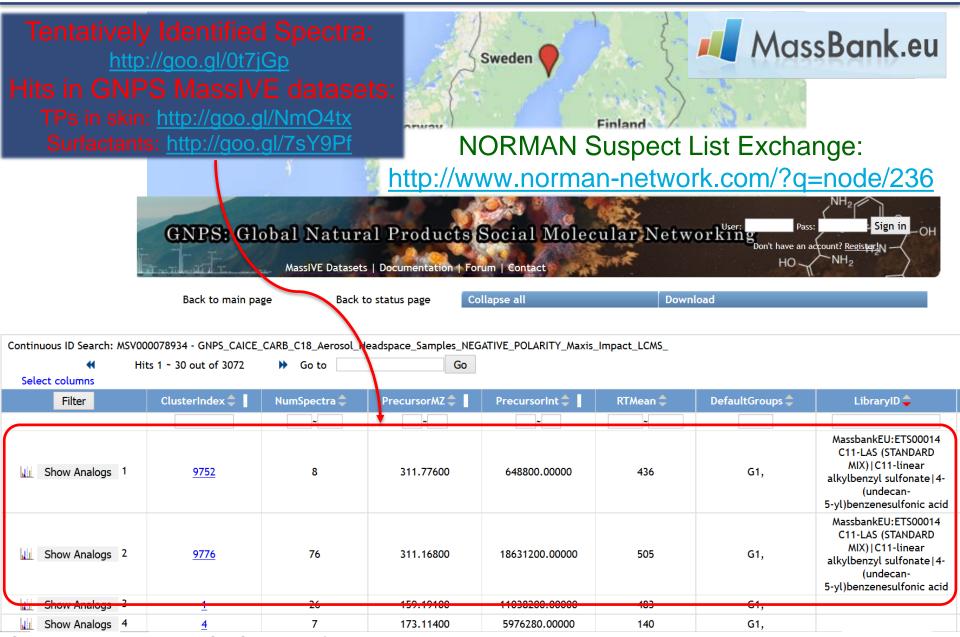
Literature MS/MS

LIT00034, LIT00037

Std Mix., Sample ETS00012, ETS00018

European (World-)Wide Exchange of Suspects





Schymanski et al. 2015, ABC, DOI: 10.1007/s00216-015-8681-7;

NORMAN Digital Sample Freezing Platform



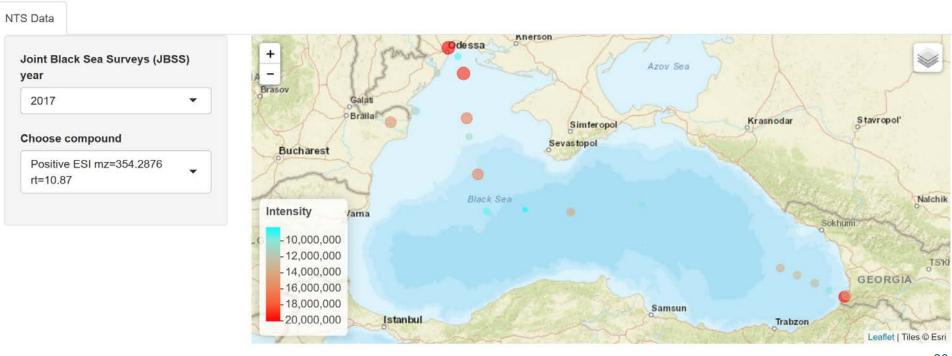
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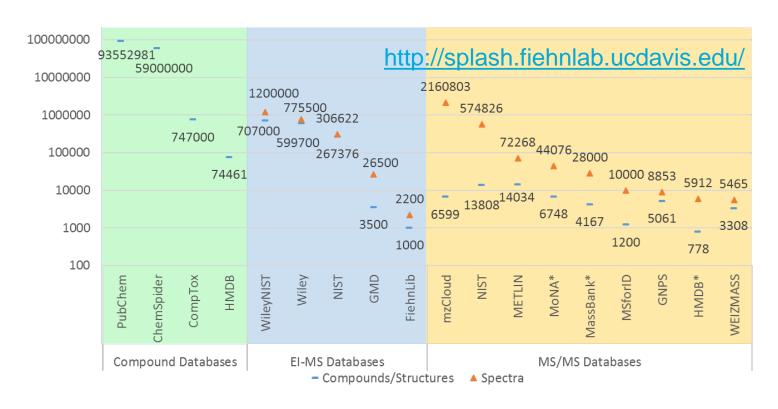
Future work: use results of **unknowns** to drive prioritization efforts

EMBLAS-II: EU/UNDP project Improving Environmental Monitoring in the Black Sea

Non-target Screening Data



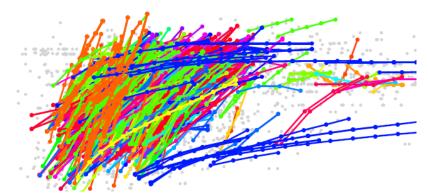
- Identifying small molecules requires information from many sources
 - Extensive compound databases available
 - Many mass spectral libraries available
 - Many excellent workflows available to collate this information
 - Community initiatives to improve communication between resources



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- Exchanging expert knowledge worldwide
 - Community efforts contribute greatly to improved cross-annotation

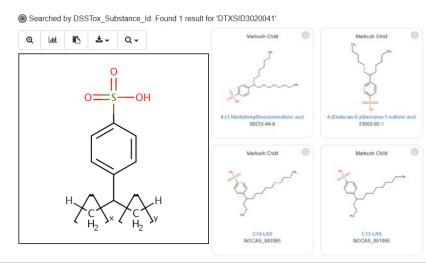


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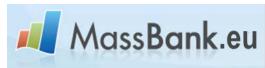


Alkylbenzenesulfonate, linear

42615-29-2 | DTXSID3020041

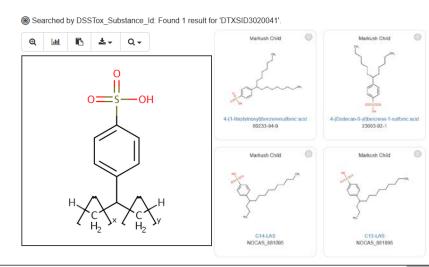


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- Tackling complex structures
 - Huge progress in cheminformatics approaches in very short time ...
 - Information in the public domain helps everyone!
 (you never know when it will help you!)



Alkylbenzenesulfonate, linear

42615-29-2 | DTXSID3020041



Acknowledgements I





















































emma.schymanski@uni.lu

Further Information:

http://www.norman-network.com/?q=node/236

https://massbank.eu/MassBank/

https://comptox.epa.gov/dashboard/

https://www.researchgate.net/project/Supporting-Mass-

Spectrometry-Through-Cheminformatics

https://github.com/MassBank/

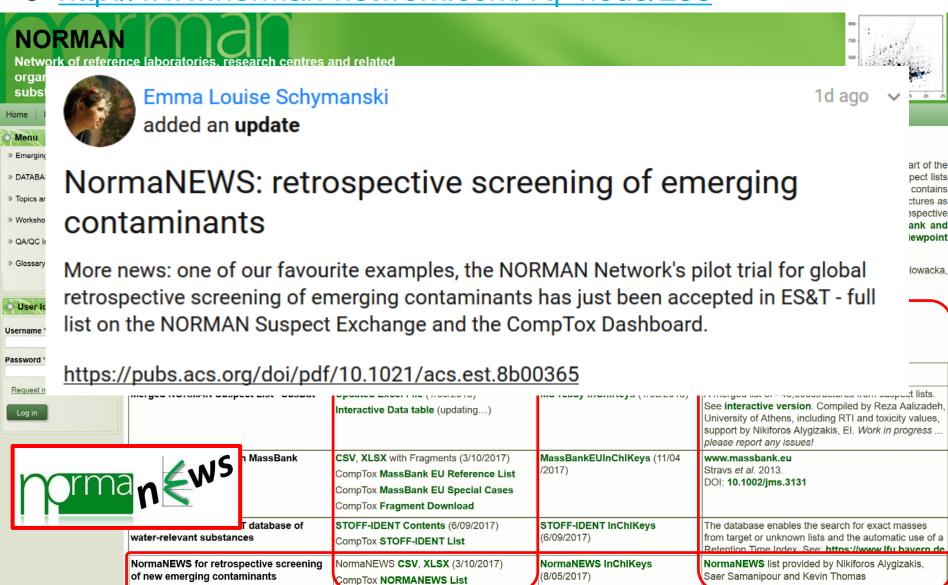




NORMAN Suspect List Exchange

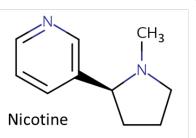


http://www.norman-network.com/?q=node/236



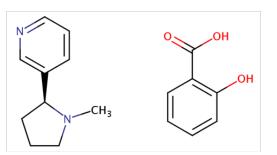
MS-Ready: Metadata & Chemical Forms





CN1CCC[C@H]1C1=CN=CC=C1 DTXSID1020930| SNICXCGAKADSCV 54-11-5 | **162.1157**| 0.929| **72**

Tox: yes | Expo: yes | Bioassay: yes



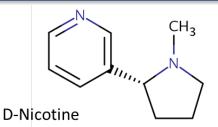
Benzoic acid, 2-hydroxy-, compd. with 3-[(2S)-1-methyl-2-pyrrolidinyl]pyridine (1:1)

OC(=O)C1=C(O)C=CC=C1.CN1CCC[C@H]1C1=CN=CC=C1

DTXSID5075319 | AIBWPBUAKCMKNS

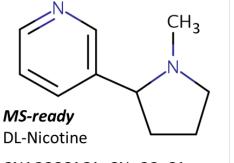
29790-52-1 | 300.1474 | 0.929 | 6

Tox: no | Expo: yes | Bioassay: no



CN1CCC[C@@H]1C1=CN=CC=C1 DTXSID0046351| SNICXCGAKADSCV 25162-00-9 | **162.1157**| 0.929| **20**

Tox: no | Expo: yes | Bioassay: yes



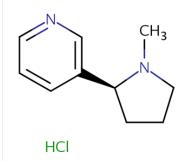
CN1CCCC1C1=CN=CC=C1

DTXSID3048154 | SNICXCGAKADSCV 22083-74-5 | **162.1157** | 0.953 | 9

Tox: yes | Expo: no | Bioassay: yes

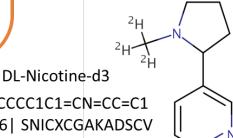
LEGEND: Name, SMILES DTXSID | InChIKey 1st Block

CAS | Monoiso. Mass | logP | Sources
Data on: Toxicity | Exposure | Bioassays



Nicotine hydrochloride

CI.CN1CCC[C@H]1C1=CN=CC=C1 DTXSID6020931| HDJBTCAJIMNXEW 2820-51-1 | **198.0924** | 0.929| **9** Tox: **no**| Expo: **yes**| Bioassay: **yes**



[2H]C([2H])([2H])N1CCCC1C1=CN=CC=C1 DTXSID80442666| SNICXCGAKADSCV 69980-24-1| **165.1345**| 0.929| **1**

Tox: no | Expo: no | Bioassay: no

MetFrag2.3: Non-target Identification



