

PubChemLite Lab: *Cheminformatics* for the *Exposome*



Assoc. Prof. Dr. Emma L. Schymanski
(plus many, many colleagues and collaborators!)

Environmental Cheminformatics Group,
Luxembourg Centre for Systems Biomedicine, University of Luxembourg

Email: emma.schymanski@uni.lu and [@ESchymanski](https://twitter.com/ESchymanski)

Web: https://www.unil.lu/lcsb/research/environmental_cheminformatics/

The Exposome Boot Camp
[Live-Stream](#), July 22-23, 2021



Slides available at
DOI: [10.5281/zenodo.5115498](https://doi.org/10.5281/zenodo.5115498)

Outline for today

- Background

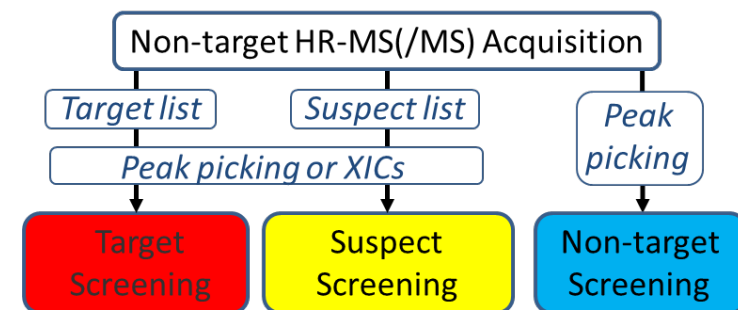
- Environmental cheminformatics & HR-MS
- Non-target screening and identification confidence
- Compound databases and spectral libraries

- Introduction to PubChemLite and MetFrag

- Examples (MassBank+PubChemLite+MetFrag)

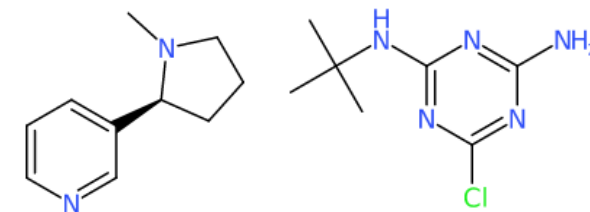
- Nicotine – relating identifications to disease outcomes
- Desethylterbutylazine – isobars and the role of metadata

- Closing – Perspectives and Community Contribution

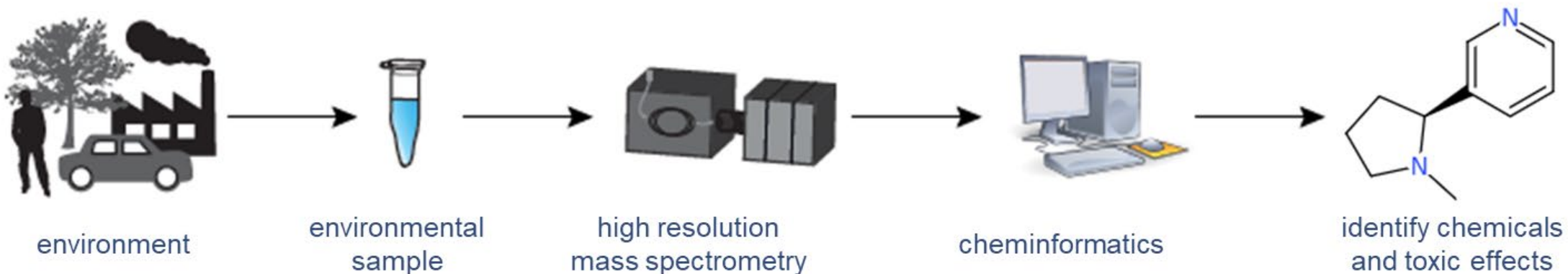


PubChemLite
EXPOSOMICS

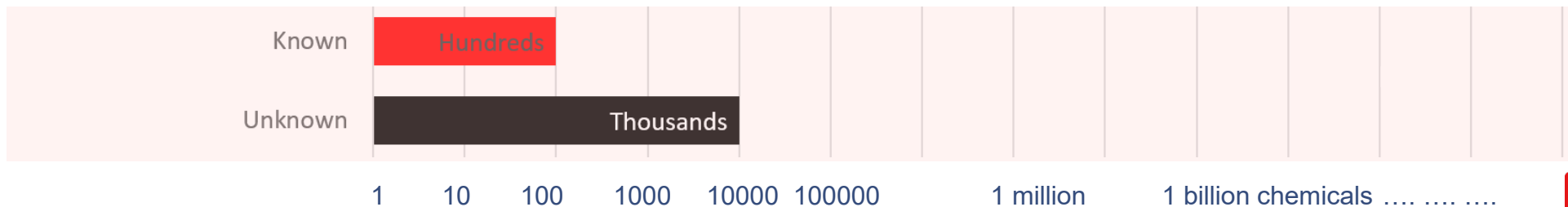
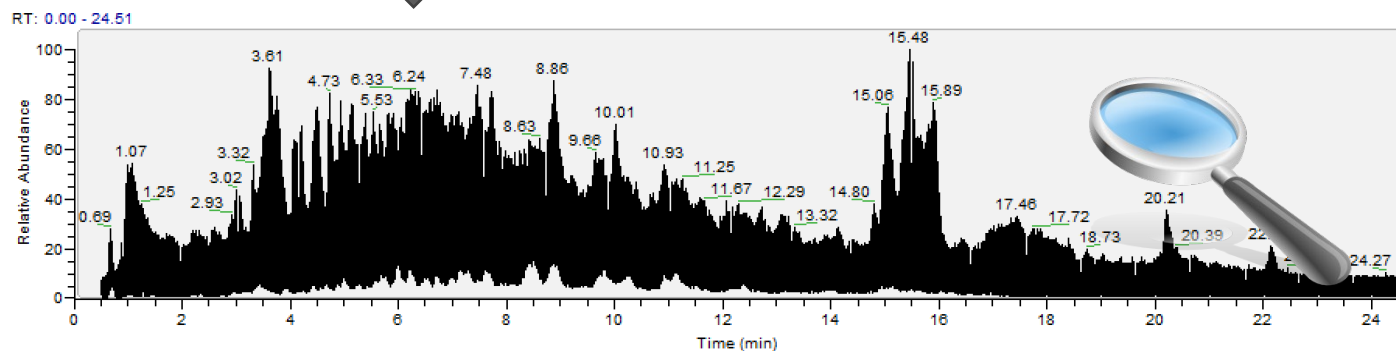
MetFrag



Background: Environmental Cheminformatics & HR-MS

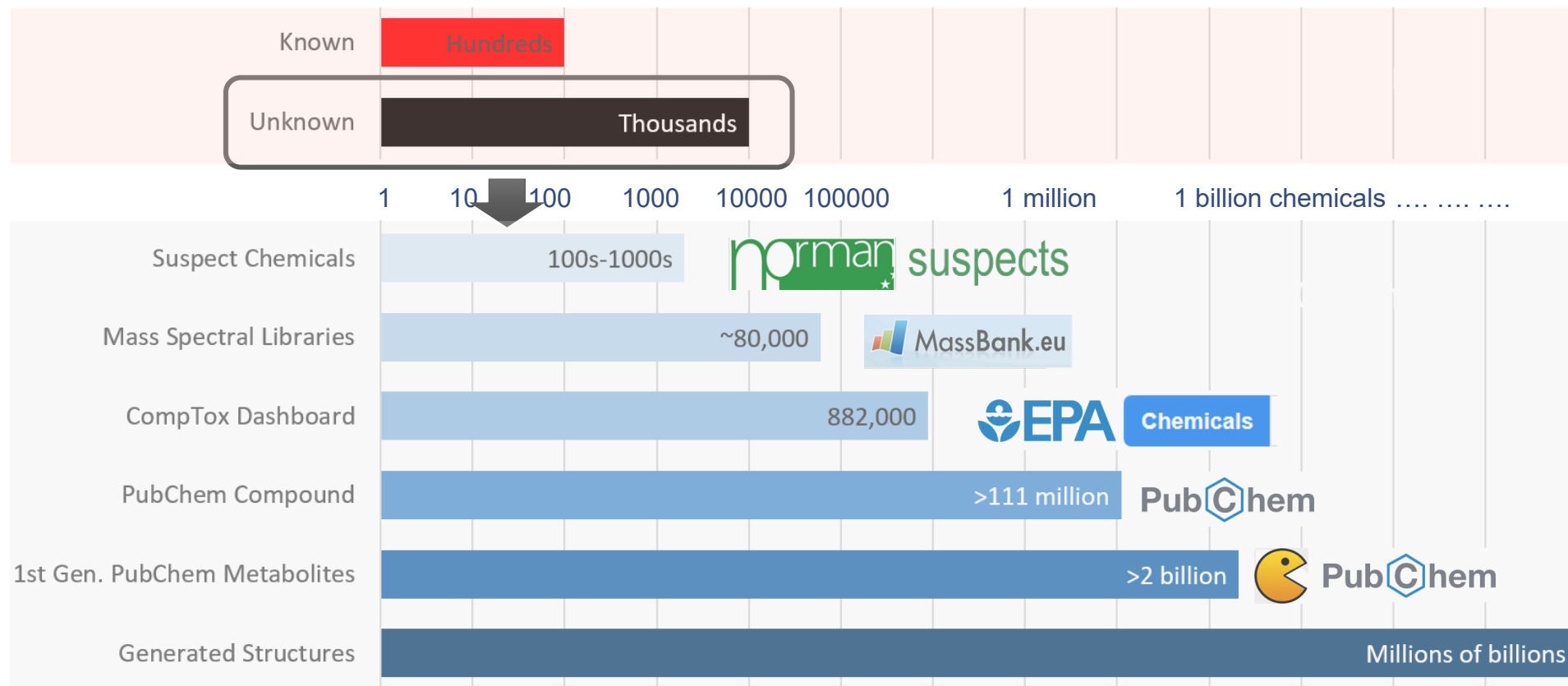
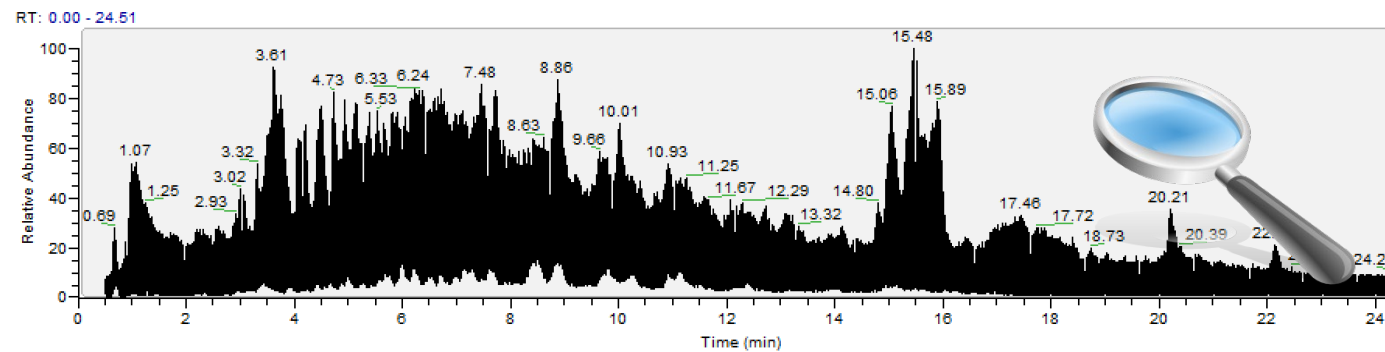


High resolution
mass spectrometry

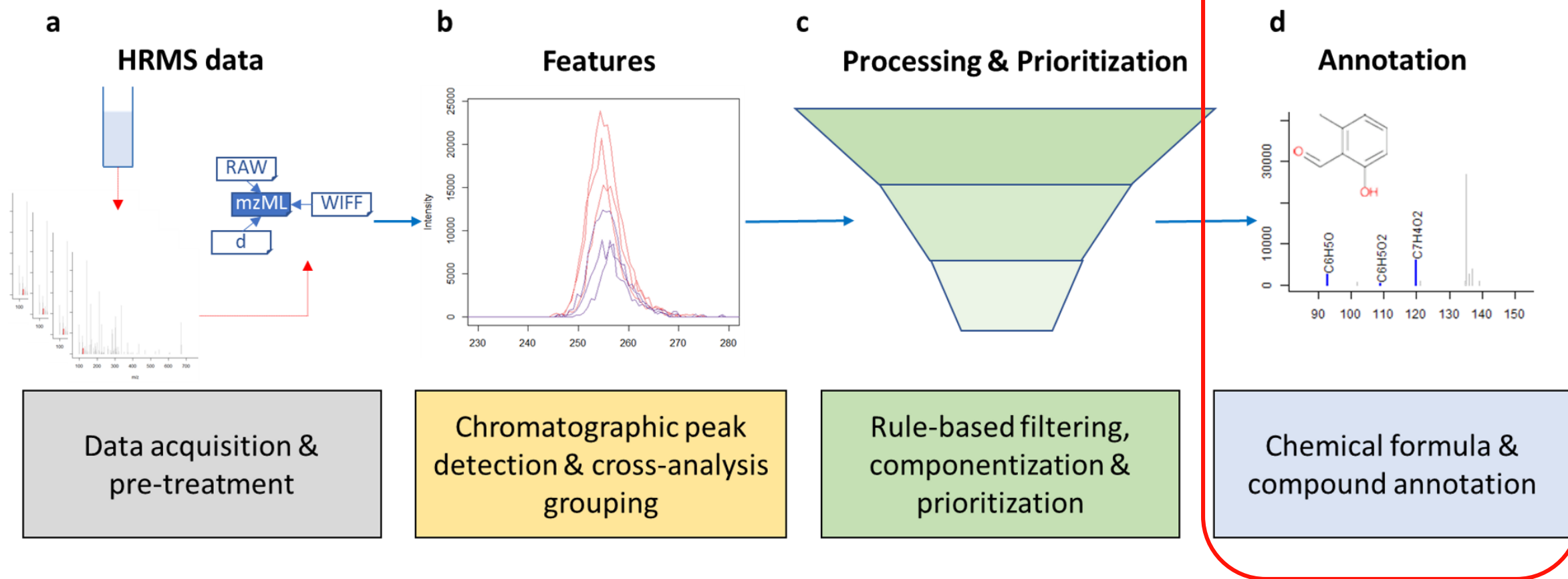


Background: Environmental Cheminformatics & HR-MS

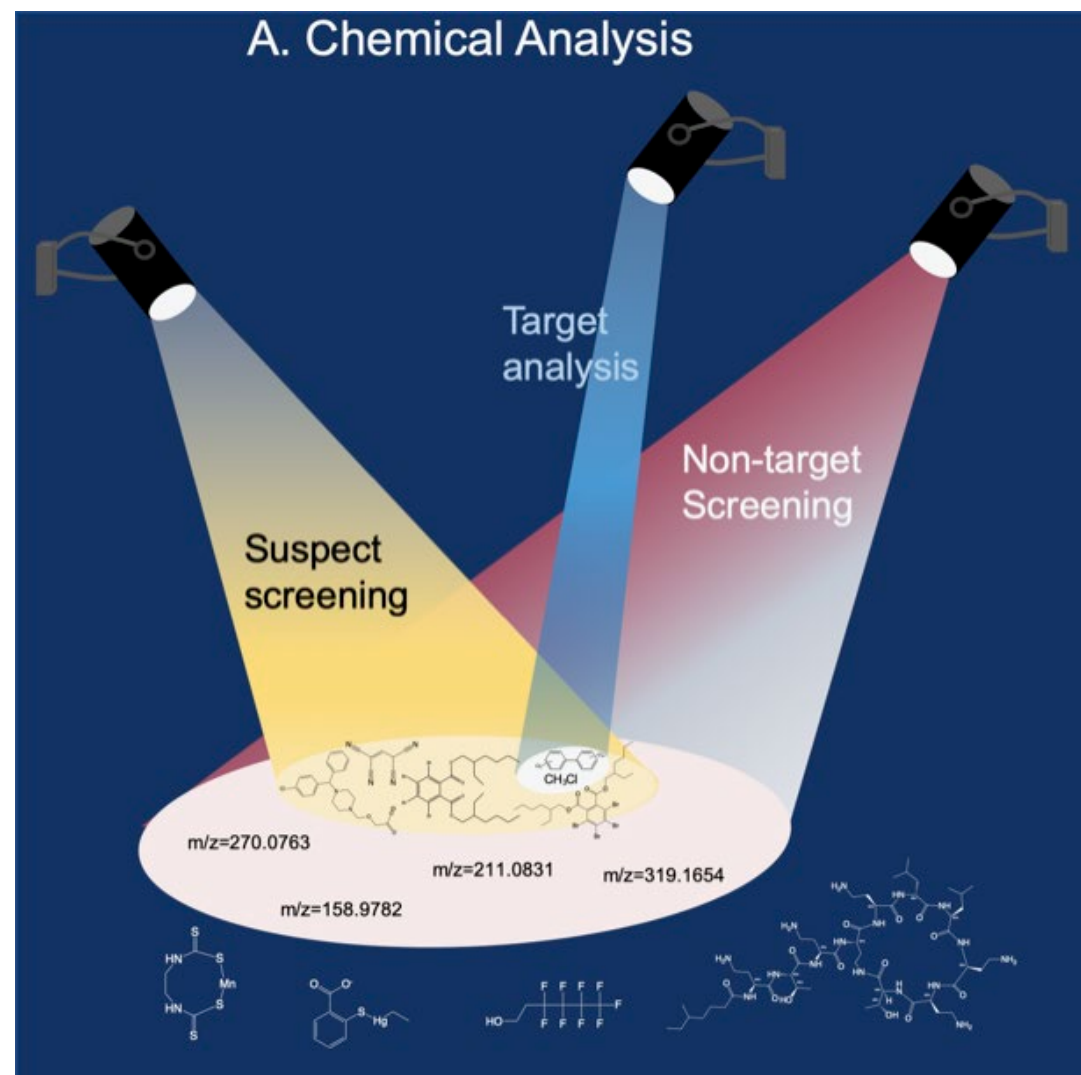
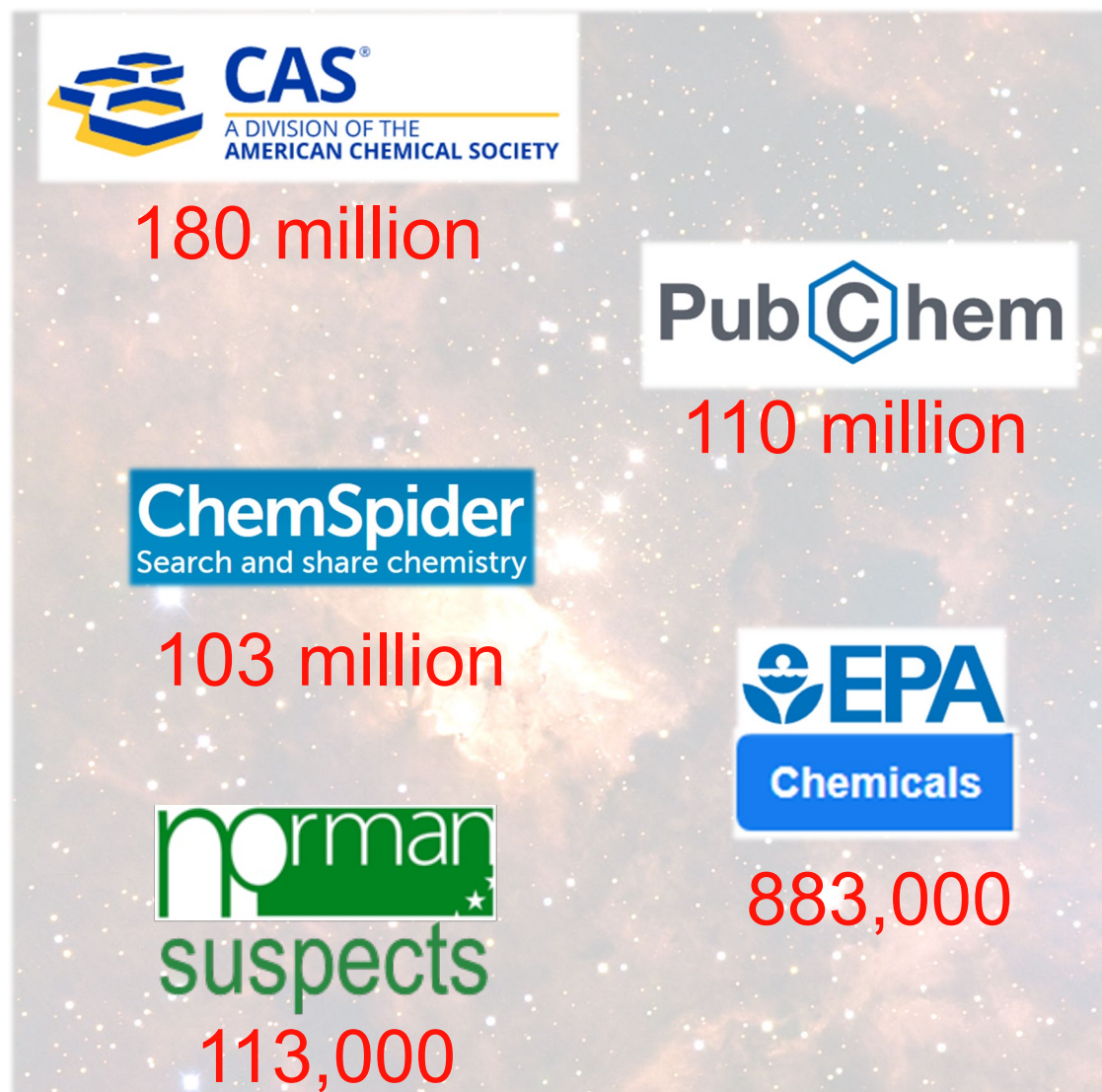
High resolution
mass spectrometry
AND connecting
chemical knowledge



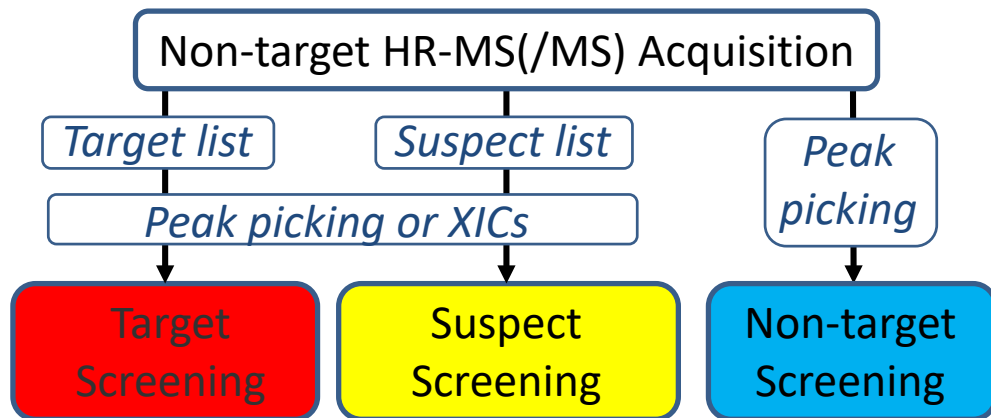
Background: Identification with HR-MS



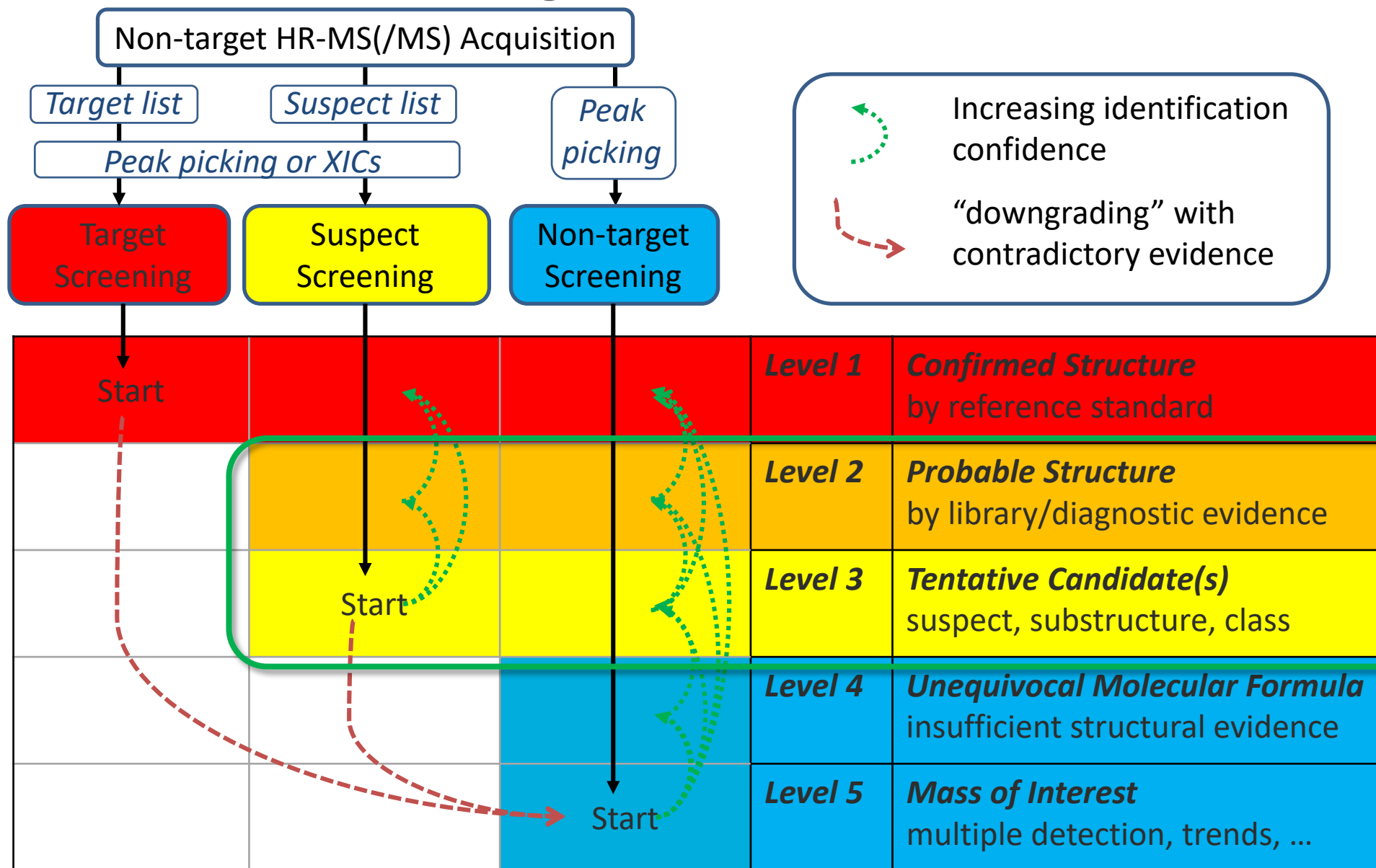
The Problem: Which chemicals are relevant? How to find them?



Identification Strategies and Confidence in NT-HRMS(/MS)



Identification Strategies and Confidence in NT-HRMS(/MS)



MassBank
High Quality Mass Spectral Database



eawag
aquatic research



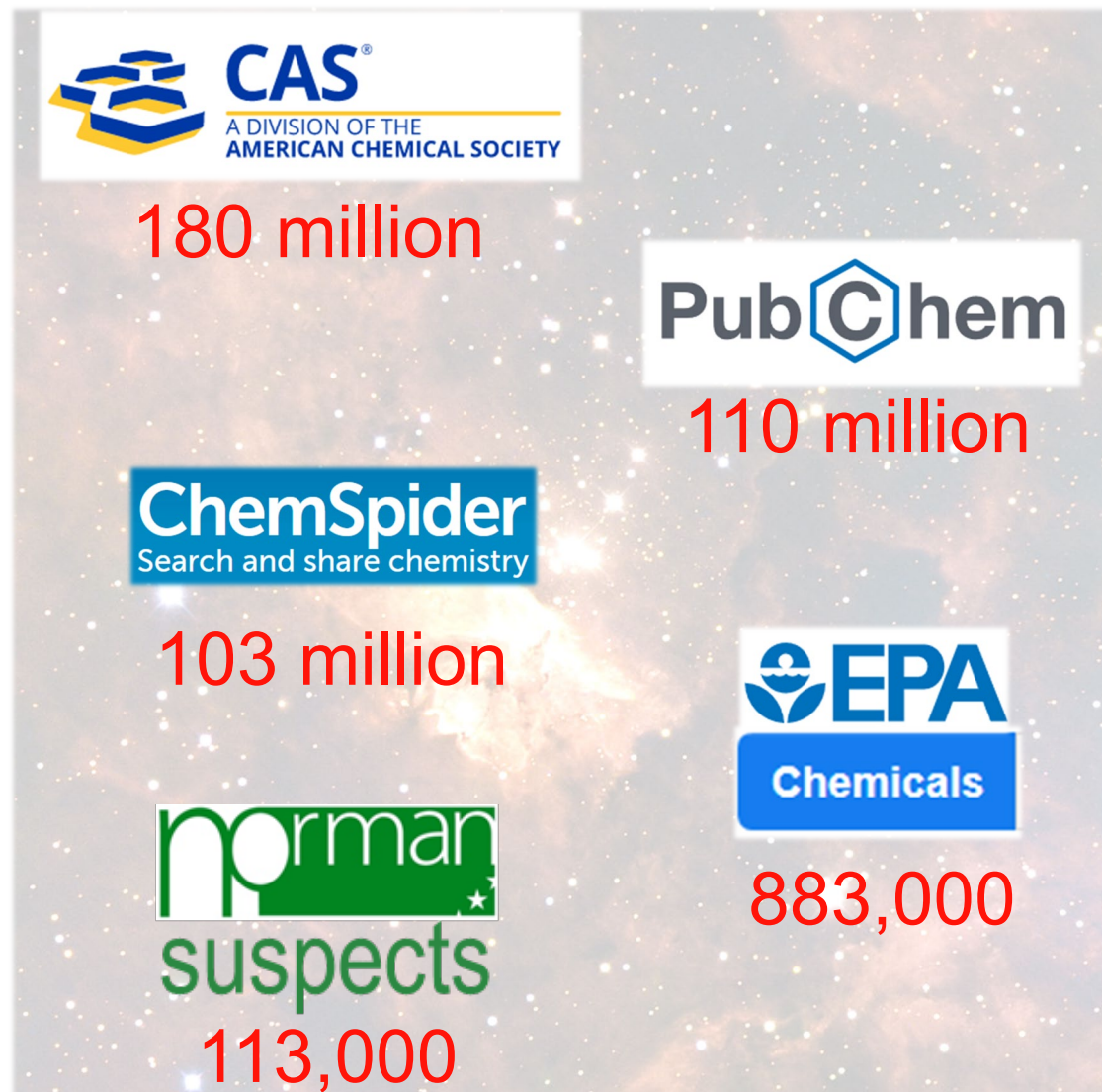
Background: Key Resources for Identification / Annotation

- Compound Databases

- A collection of structures, their properties and associated information
- Generally little or no spectral data, rather structures & links
- Largest have >100 million structures in them
 - But don't assume that everything is in there – it isn't!!!!



370,000 entries “small”



Background: Key Resources for Identification / Annotation

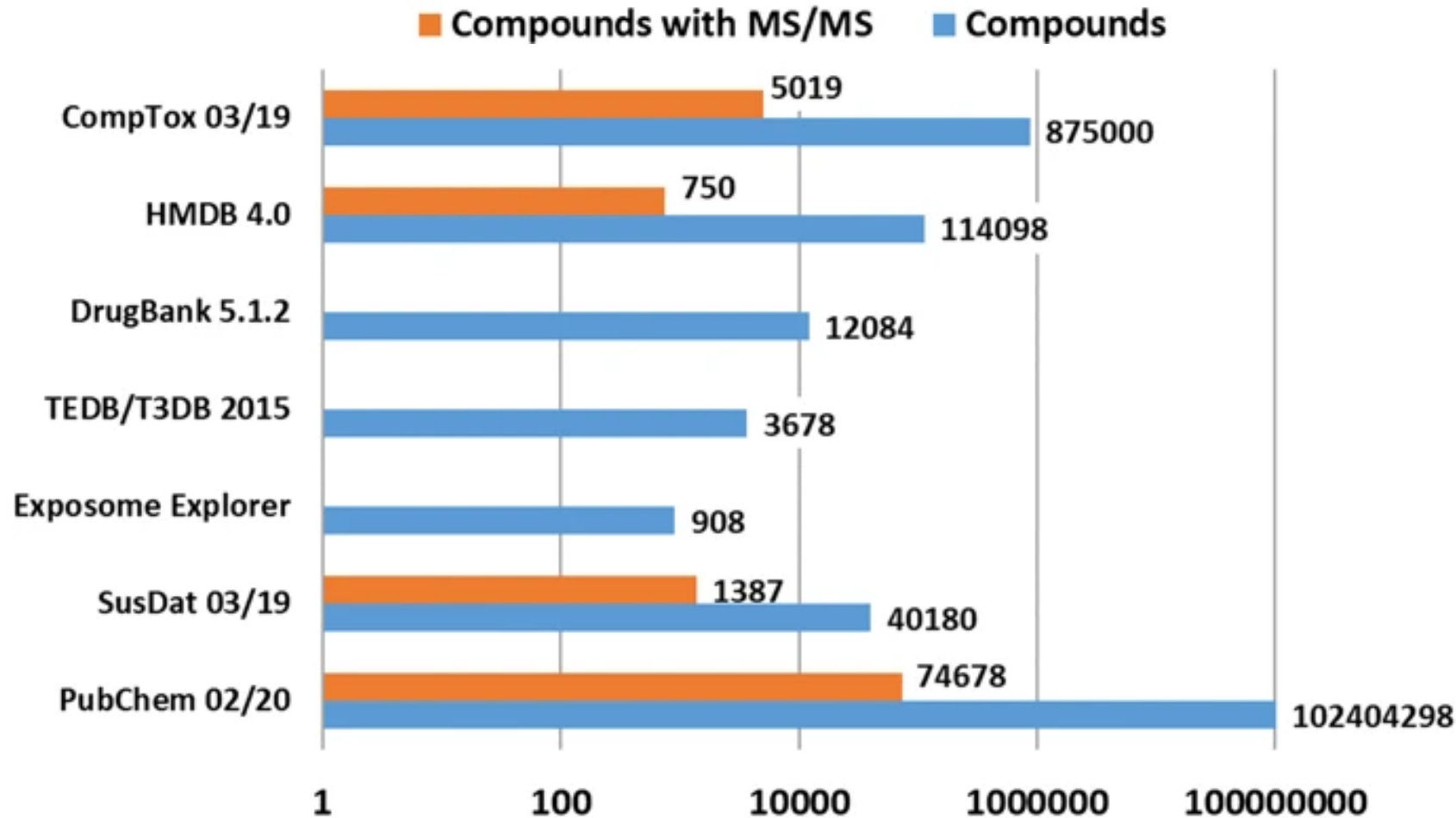
- Mass Spectral Databases or Libraries
 - A collection of structures, mass spectra and associated information
 - NIST and Wiley are widely accepted for GC-EI/MS
 - Together >1.2 million spectra of 707,000 compounds
- MS/MS databases are growing, none are yet “established”
 - Together > 2 million spectra, but only approx. 40-80,000 compounds
 - MS/MS available for only ~0.1-4 % of relevant exposomics resources (see next slide)
- Key resource(s) mentioned today:
 - MassBank EU: <http://massbank.eu/MassBank>
 - MoNA: <https://mona.fiehnlab.ucdavis.edu/>

 **MassBank**
High Quality Mass Spectral Database

MoNA
MassBank of North America

Background: Scarcity of MS/MS Spectra in Exposomics

- MS/MS avail. for ~0.1-4 % of relevant exposomics resources

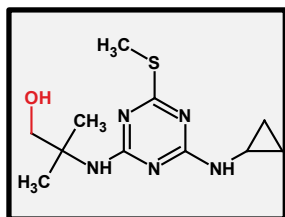


Identification Strategies and Confidence in NT-HRMS(/MS)

Example

Identification confidence

Minimum data requirements



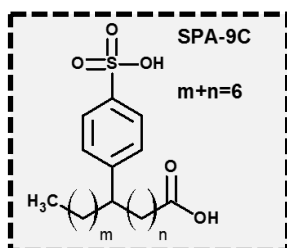
Level 1: Confirmed structure
by reference standard

MS, MS², RT, **Reference Std.**

Level 2: Probable structure
a) by library spectrum match
b) by diagnostic evidence

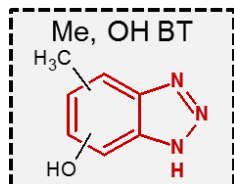
MS, MS², **Library MS² >0.9 sim**
MS, MS², Exp. data

 **MassBank**
High Quality Mass Spectral Database

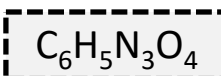


Level 3: Tentative candidate(s)
structure, substituent, class

MS, MS², Exp. data



 **Metrag**



Level 4: Unequivocal molecular formula

MS isotope/adduct

192.0757

Level 5: Exact mass of interest

MS

 **mini**

 **LCSB**

Outline for today

- Background

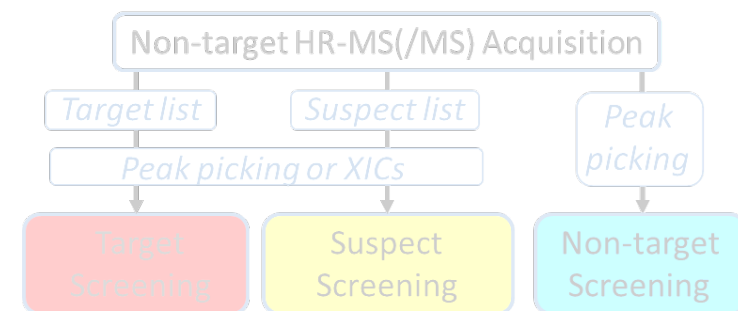
- Environmental cheminformatics & HR-MS
- Non-target screening and identification confidence
- Compound databases and spectral libraries

- Introduction to PubChemLite and MetFrag

- Examples (MassBank+PubChemLite+MetFrag)

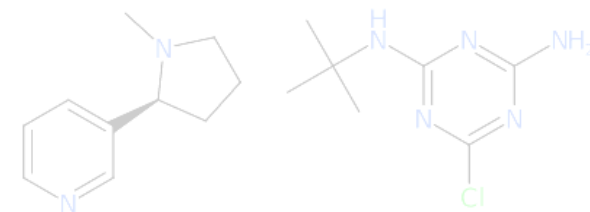
- Nicotine – relating identifications to disease outcomes
- Desethylterbutylazine – isobars and the role of metadata

- Closing – Perspectives and Community Contribution



PubChemLite
EXPOSOMICS

MetFrag



Introduction to MetFrag

<https://msbi.ipb-halle.de/MetFrag/>

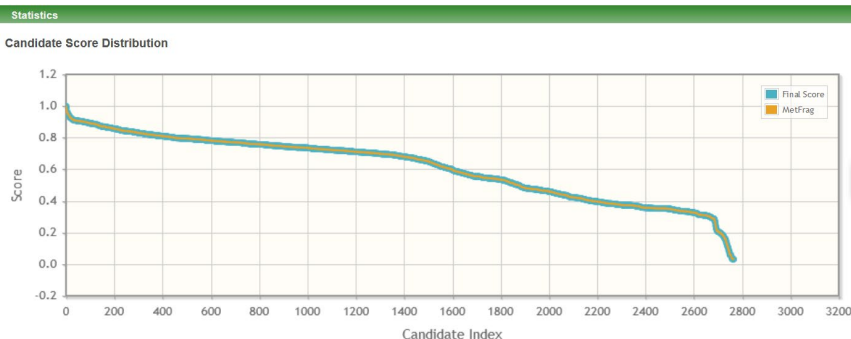
m/z $[M-H]^-$
213.9637
 ± 5 ppm

5 ppm
0.001 Da

PubChem



Ranked Candidates



MS/MS

134.0054	339689
150.0001	77271
213.9607	632466



Key Challenge: MS and MS/MS alone is not enough!



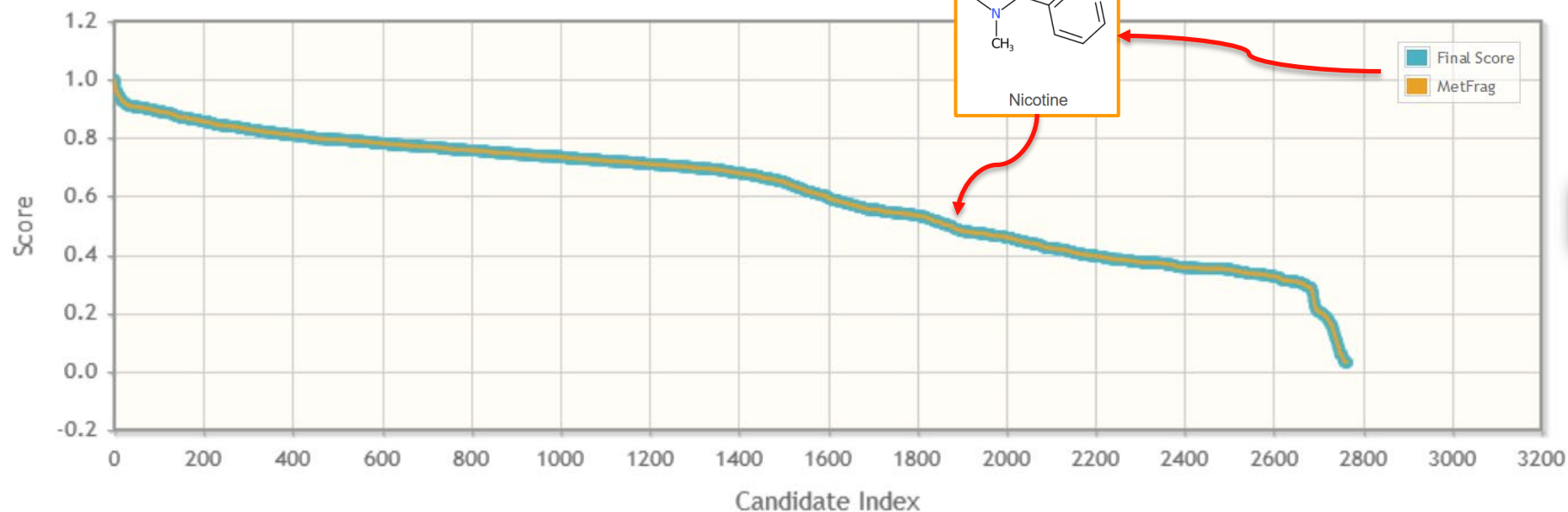
Key Challenge: MS and MS/MS alone is not enough!



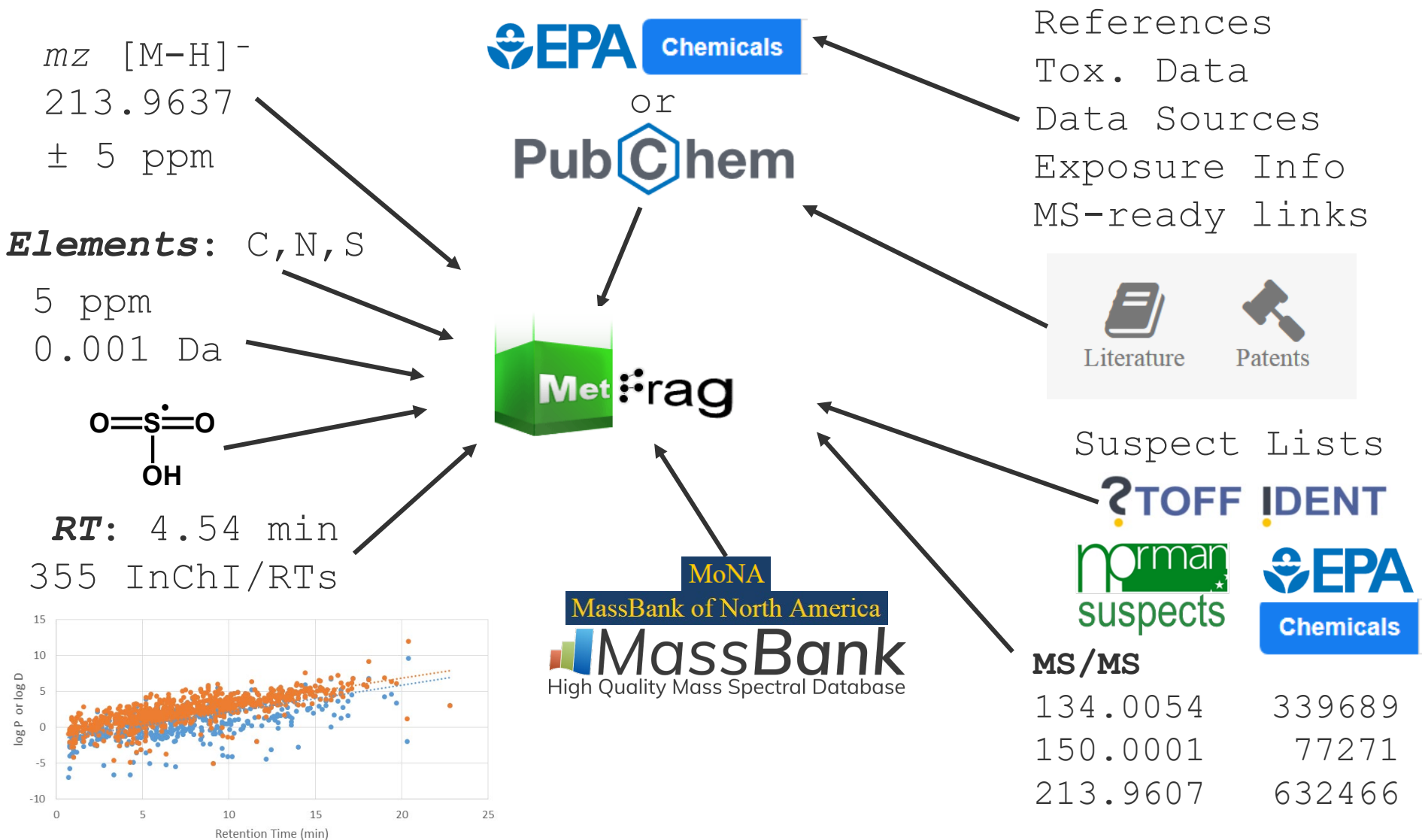
PubChem

Statistics

Candidate Score Distribution



Status Quo in 2016: MetFrag Relunched ...



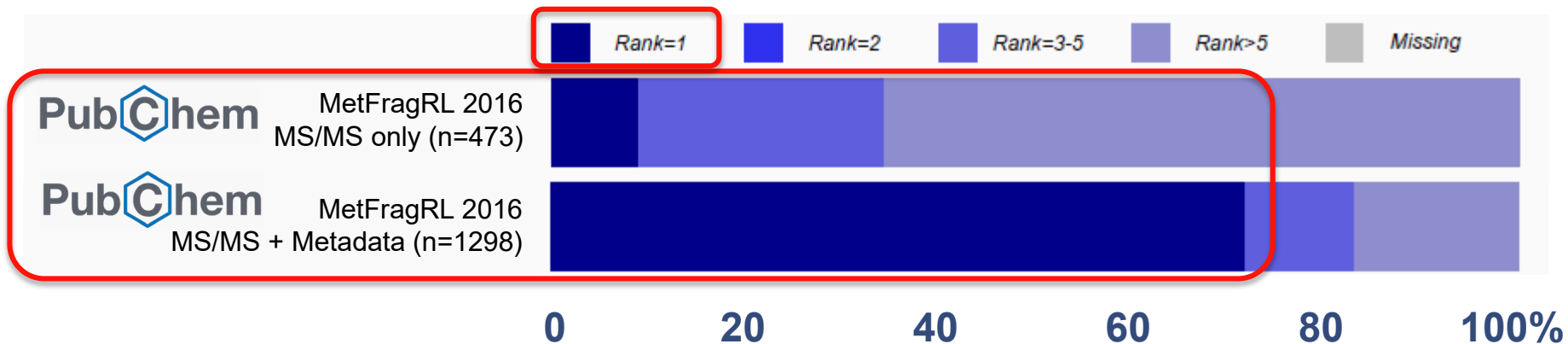
eawag
aquatic research

solutions



MetFragRL + PubChem + MS/MS + Metadata

- Adding literature, references & RT boosts to ~71 % rank 1!



MetFragRL + PubChem + MS/MS + Metadata



BUT ...databases grow ... ID performance drops

... and run times rise ... (a lot!)



Problem: Exposomics “Chemical Space” is too big!



180 million



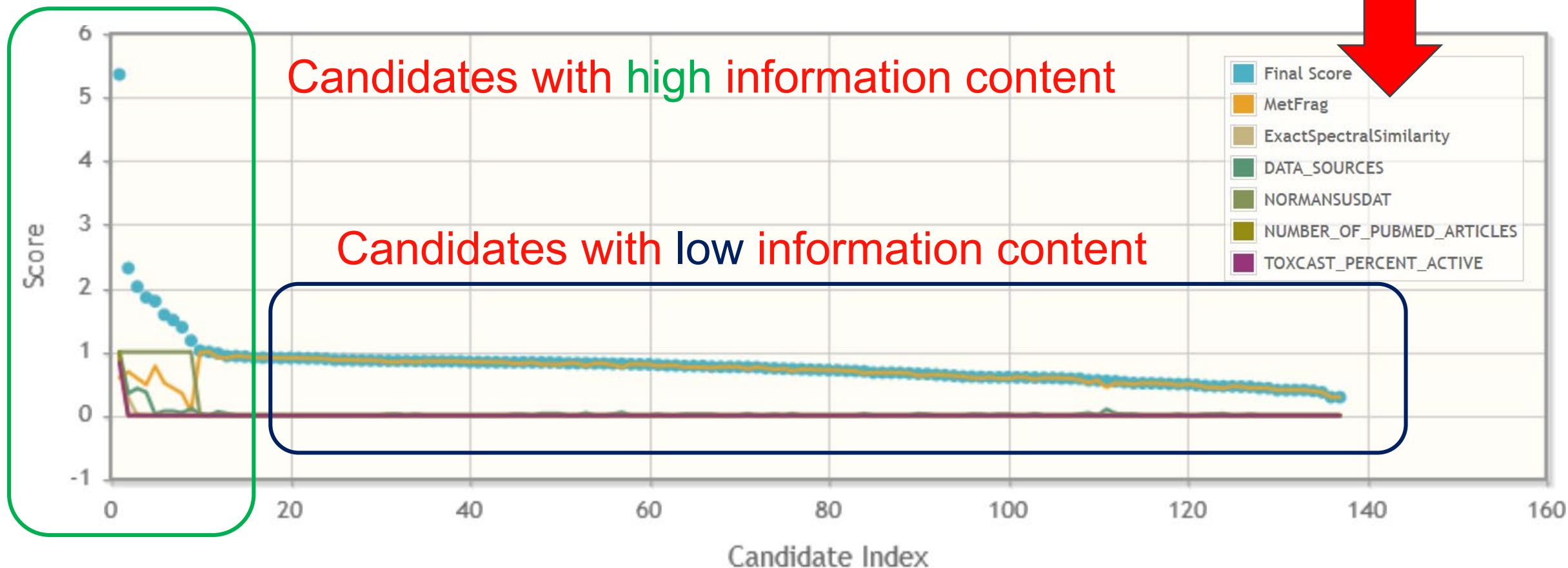
110 million



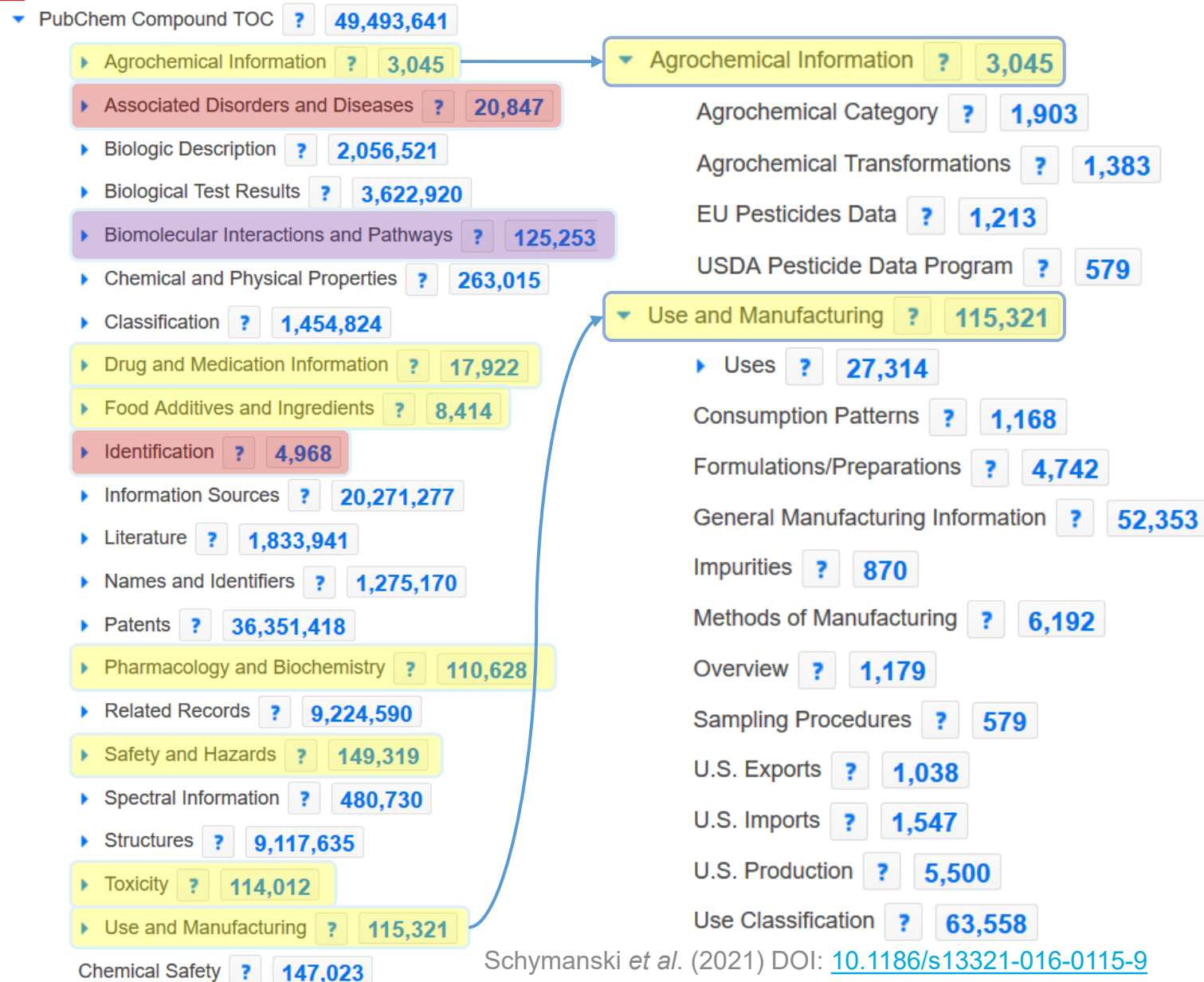
103 million



883,000



Can we break down PubChem into useful bits?



PubChem Furathiocarb (Compound)

CONTENTS



7 Agrochemical Information



7.1 Agrochemical Category



Insecticides

▶ EU Pesticides Database

7.2 Agrochemical Transformations



Furathiocarb has known environmental transformation products that include [carbofuran](#).

S60 | SWISSPEST19 | Swiss Pesticides and Metabolites from Kiefer et al 2019 | DOI:10.5281/zenodo.3544759

▶ NORMAN Suspect List Exchange

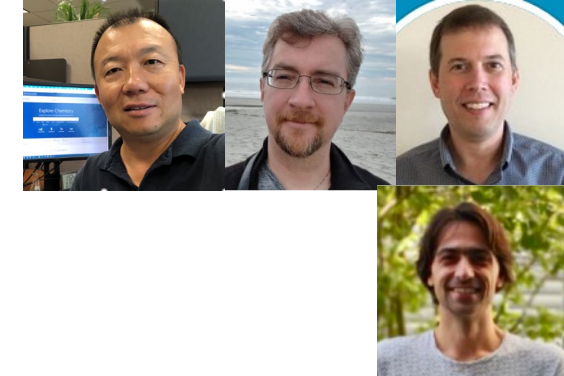
7.3 EU Pesticides Data



Active Substance	furathiocarb
Status	Not Approved [Reg. (EC) No 1107/2009]
Categories	Insecticides

<https://pubchem.ncbi.nlm.nih.gov/compound/Furathiocarb#section=Agrochemical-Information>

Introducing ...



▼ PubChem Compound TOC ? 49,493,641

▶ Agrochemical Information ? 3,045

▶ Associated Disorders and Diseases ? 20,847

▶ Biologic Description ? 2,056,521

▶ Biological Test Results ? 3,622,920

▶ Biomolecular Interactions and Pathways ? 125,253

▶ Chemical and Physical Properties ? 263,015

▶ Classification ? 1,454,824

▶ Drug and Medication Information ? 17,922

▶ Food Additives and Ingredients ? 8,414

▶ Identification ? 4,968

▶ Information Sources ? 20,271,277

▶ Literature ? 1,833,941

▶ Names and Identifiers ? 1,275,170

▶ Patents ? 36,351,418

▶ Pharmacology and Biochemistry ? 110,628

▶ Related Records ? 9,224,590

▶ Safety and Hazards ? 149,319

▶ Spectral Information ? 480,730

▶ Structures ? 9,117,635

▶ Toxicity ? 114,012

▶ Use and Manufacturing ? 115,321

Chemical Safety ? 147,023

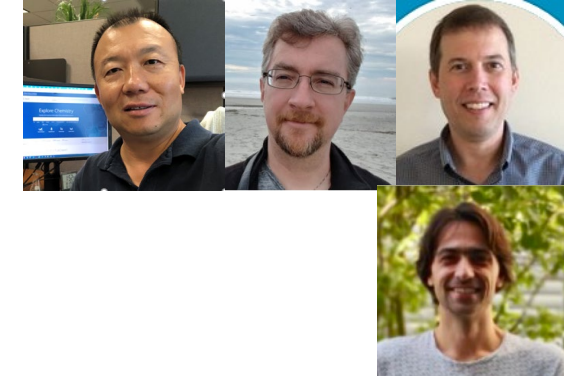
PubChemLite
EXPOSOMICS

~370,000 entries “small”

Schymanski *et al.* (2021) DOI: [10.1186/s13321-016-0115-9](https://doi.org/10.1186/s13321-016-0115-9)

Introducing ...

PubChemLite EXPOSOMICS



Collapsed by InChIKey First Block (skeleton)
and by presence of annotation content

~370,000 entries “small”

PubChem

Compounds (6) Substances (2)

Searching chemical names and synonyms including IUPAC names and InChIKeys across the compound collection. Note that annotations text searched. [Read More...](#)

6 results SORT BY



2,3-dihydroxy-2,3-dihydrobenzoic Acid; 5,6-dihydroxycyclohexa-1,3-diene-1-carboxylic acid; 100459-00-5; 2,3-dihydro-2,3-dihydroxybenzoic Acid; ACM

Compound CID (sort by): 3

MF: C₇H₈O₄ MW: 156.14g/mol

InChIKey: [INCSWKICYAHB-UHFFFAOYSA-N](#)

IUPAC Name: 5,6-dihydroxycyclohexa-1,3-diene-1-carboxylic acid

Create Date: 2004-09-16

[Summary](#) [Similar Structures Search](#) [Related Records](#)



(2S,3S)-2,3-dihydroxy-2,3-dihydrobenzoic Acid; 176487-06-2; (2S,3S)-2,3-dihydroxybenzoate; (5S,6S)-5,6-dihydroxycyclohexa-1,3-diene-1-carboxylic acid; (2S,3S)-2,3-Dihydro-2,3-dihydroxybenzoate; ...

Compound CID (sort by): 9964159

MF: C₇H₈O₄ MW: 156.14g/mol

InChIKey: [INCSWKICYAHB-WDSKDSINSA-N](#)

IUPAC Name: (5S,6S)-5,6-dihydroxycyclohexa-1,3-diene-1-carboxylic acid

Create Date: 2006-10-25

zenodo [Communities](#)

October 31, 2020

[Dataset](#) [Open Access](#)

[Edit](#)

[New version](#)

PubChemLite for Exposomics

[Bolton, Evan](#); [Schymanski, Emma](#); [Kondic, Todor](#); [Thiessen, Paul](#); [Zhang, Jeff](#)

PubChemLite is a subset of PubChem (<https://pubchem.ncbi.nlm.nih.gov/>) selected from major contents page at the PubChem Classification Browser (<https://pubchem.ncbi.nlm.nih.gov/classification/#hid=72>). With this release, there is now just one is the former tier1 plus two new categories (Associated Disorders &

is 371,663 compounds (31 Oct 2020) compiled from 10 categories: DrugMedicInfo, FoodRelated, Pharmacolnfo, Info, Identification.

Collapsed by InChIKey first block, reporting the most CIDs. Entries that will be ignored by MetFrag (e.g. transition metals) have been removed on the PubChem FTP site. The "AnnoTy how represented, the subsequent column (name per category) counts the ories available in the next sub-category of the TOC entry.



Met:rag

Database Settings

Database:

Neutral Mass: Search ppm:

Formula:

Identifiers:

[Retrieve Candidates](#) [4 Candidates](#)

Communities

LCSB Environmental Cheminformatics Group [Remove](#)

1,084 views 1,170 downloads
[See more details...](#)

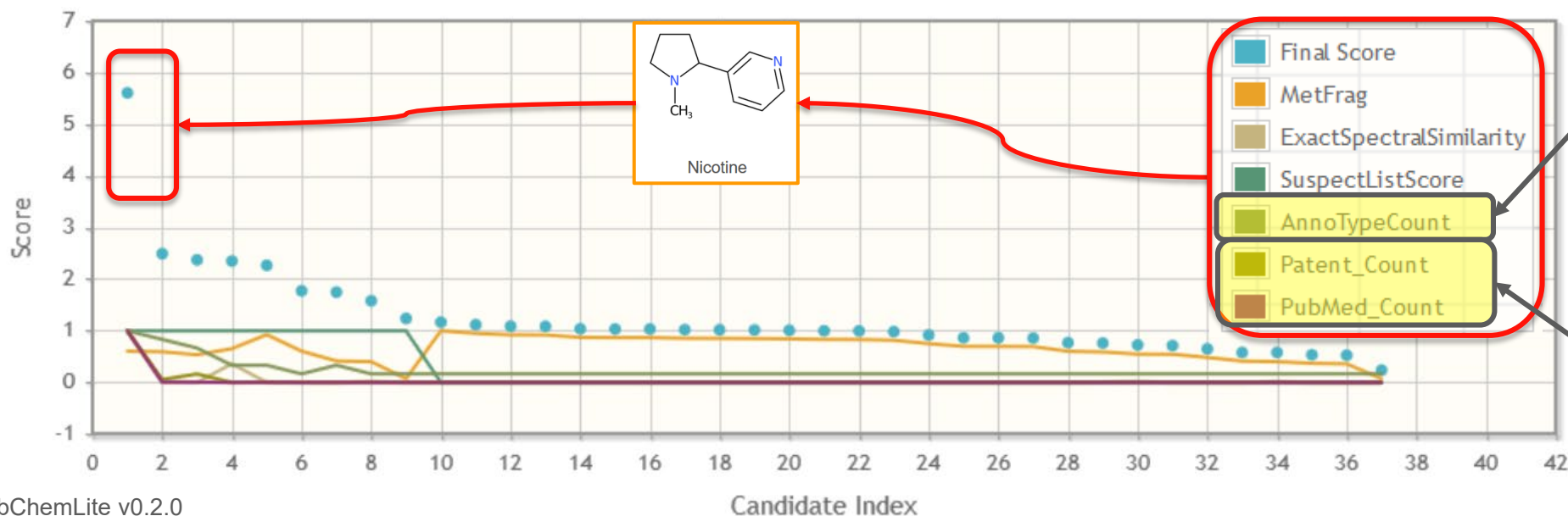
Schymanski *et al.* (2021) DOI: [10.1186/s13321-016-0115-9](https://doi.org/10.1186/s13321-016-0115-9)

MetFragRL + PubChemLite: tailor-made database + metadata



Statistics

Candidate Score Distribution



PubChem Compound TOC	49,493,641
Agrochemical Information	3,045
Associated Disorders and Diseases	20,847
Biologic Description	2,056,521
Biological Test Results	3,622,920
Biomolecular Interactions and Pathways	125,253
Chemical and Physical Properties	263,015
Classification	1,454,824
Drug and Medication Information	17,922
Food Additives and Ingredients	8,414
Identification	4,968
Information Sources	20,271,277
Literature	1,833,941
Names and Identifiers	1,275,170
Patents	36,351,418
Pharmacology and Biochemistry	110,628
Related Records	9,224,590
Safety and Hazards	149,319
Spectral Information	480,730
Structures	9,117,635
Toxicity	114,012
Use and Manufacturing	115,321
Chemical Safety	147,023



Literature



Patents

PubChemLite v0.2.0

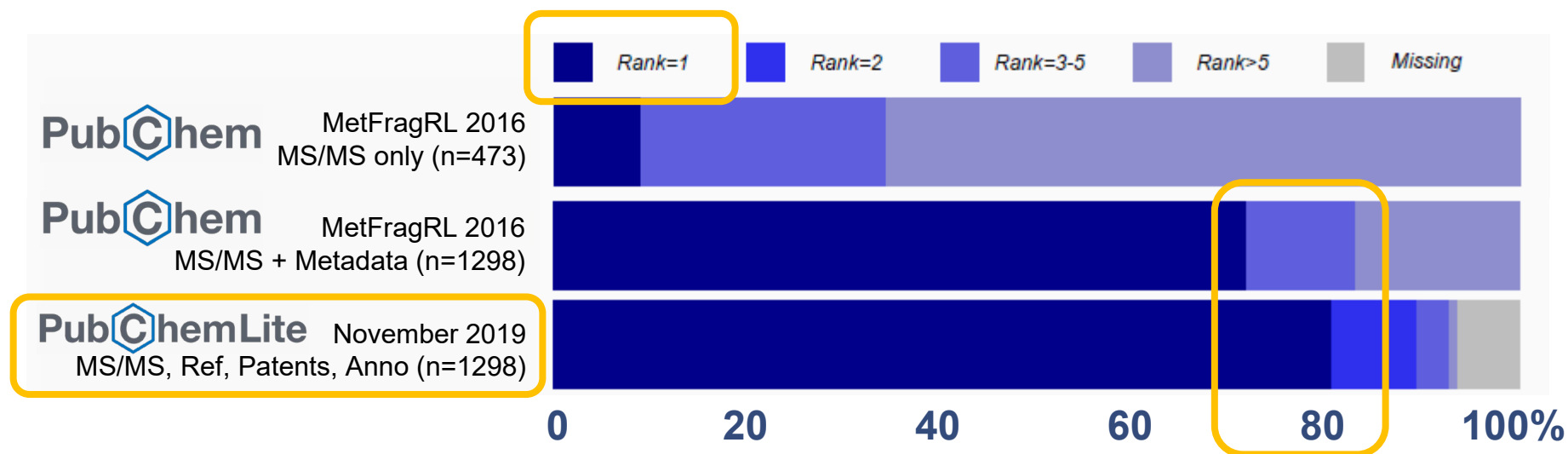
Candidate Index

MetFrag+PubChemLite+Formula+MoNA+SusDat+Pat+Refs+Anno + <https://massbank.eu/MassBank/RecordDisplay?id=EQ300804>



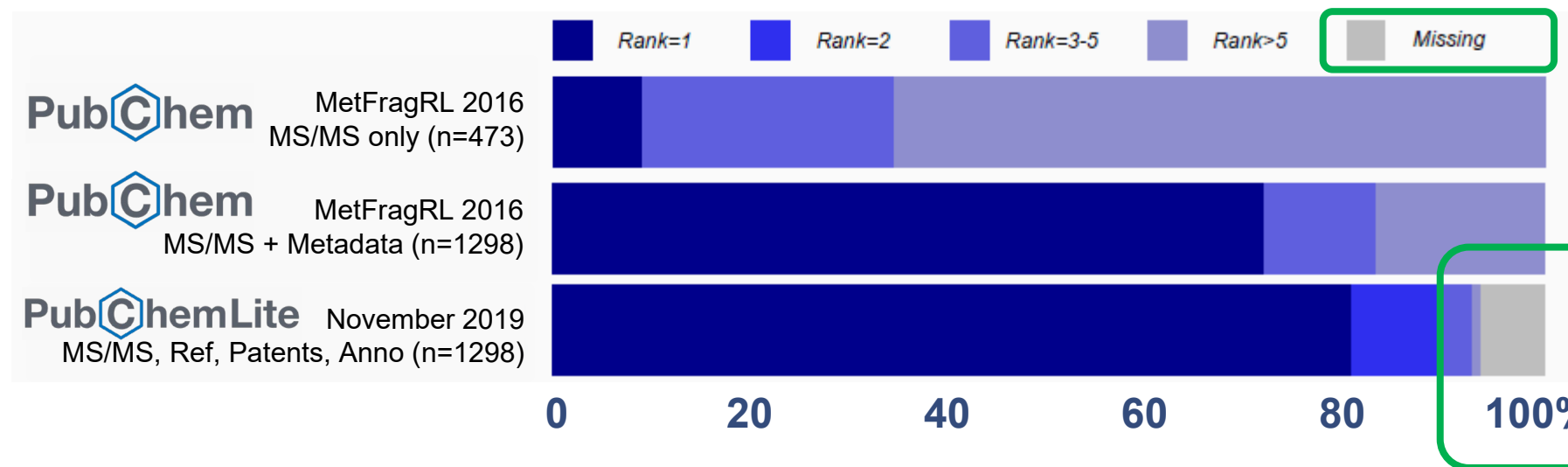
How does PubChemLite perform?

- ~110 M => ~370 K ... how does this influence performance?



How does PubChemLite perform?

- ~110 M => ~370 K ... how does this influence performance?

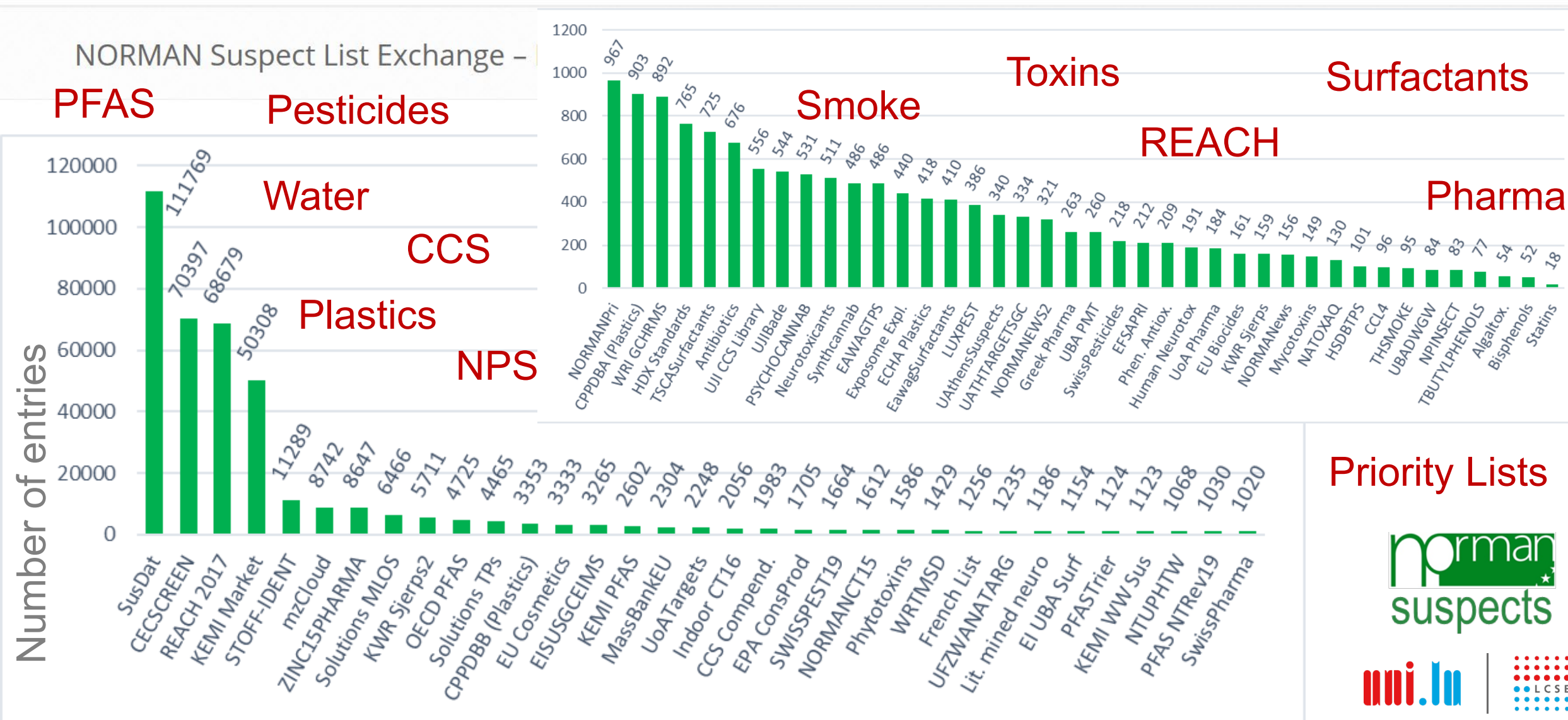


norman
suspects

Expert Knowledge: NORMAN Suspect List Exchange (>80 lists!)

<https://www.norman-network.com/nds/SLE/>

<https://zenodo.org/communities/norman-sle>



Filling Gaps: Integrating NORMAN-SLE

<https://www.norman-network.com/nds/SLE/> => <https://pubchem.ncbi.nlm.nih.gov/source/23819>

PubChem NORMAN Suspect List Exchange



Organization	NORMAN Network (c/o UniLu)
Category	Research and Development
URL	https://www.norman-network.com/nds/SLE/
License Note	Data: CC-BY 4.0; Code (hosted by ECI, LCSB): Artistic-2.0
License URL	https://creativecommons.org/licenses/by/4.0/
Contact Name	Emma Schymanski
Address	6 avenue du Swing, Belvaux, Luxembourg, 4367
Data Source ID	23819
Data in PubChem	114,573 Live Substances 16,423 Annotations 1 Classification
Last Updated	2021/04/15



Filling Gaps: Integrating NORMAN-SLE

NORMAN-SLE Classification: <https://pubchem.ncbi.nlm.nih.gov/classification/#hid=101>

PubChem Classification Browser

Help

Browse PubChem data using a classification of interest, or search for PubChem records annotated with the desired classification/term (e.g., MeSH: phenylpropionates, or Gene Ontology: DNA repair). [More...](#)

Select classification

Search selected classification by

NORMAN Suspect List Exchange

Keyword

Enter desired search term

Search

Classification description (from NORMAN Suspect List Exchange)

The NORMAN Suspect List Exchange (NORMAN-SLE) is a central access point for NORMAN members (and others) to find suspect lists relevant for their environmental monitoring questions. [More...](#)

▼ NORMAN Suspect List Exchange Classification ? ↗ 113,080

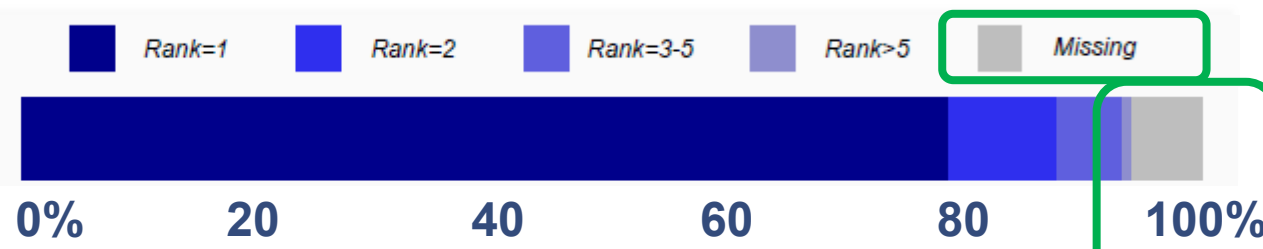
- ▶ S13 | EUCOSMETICS | Combined Inventory of Ingredients Employed in Cosmetic Products (2000) and Revised Inventory (2006) ? 3,859
- ▶ S25 | OECDPFAS | List of PFAS from the OECD ? 3,677
- ▶ S36 | UBAPMT | Potential Persistent, Mobile and Toxic (PMT) substances ? 254
- ▶ S50 | CCSCOMPEND | The Unified Collision Cross Section (CCS) Compendium ? 885
- ▶ S60 | SWISSPEST19 | Swiss Pesticides and Metabolites from Kiefer et al 2019 ? 1,343
- ▶ S61 | UJICCSLIB | Collision Cross Section (CCS) Library from UJI ? 574
- ▶ S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag ? 258
- ▶ S68 | HSDBTPS | Transformation Products Extracted from HSDB Content in PubChem ? 102
- ▶ S69 | LUXPEST | Pesticide Screening List for Luxembourg ? 386
- ▶ S72 | NTUPHTW | Pharmaceutically Active Substances from National Taiwan University ? 1,068
- ▶ S75 | CyanoMetDB | Comprehensive database of secondary metabolites from cyanobacteria ? 2,088

norman
suspects



Assessing the Missing Entries in PubChemLite

PubChemLite 14 Jan 2020
MS/MS, Ref, Patents, Anno (n=977)



- ▼ NORMAN Suspect List Exchange Classification ? ↗ 117,037
 - ▶ S13 | EUCOSMETICS | Combined Inventory of Ingredients Employed in Cosmetic Products (2000) and
 - ▶ S25 | OECDPFAS | List of PFAS from the OECD ? 3,680
 - ▶ S50 | CCSCOMPEND | The Unified Collision Cross Section (CCS) Compendium ? 647
 - ▶ S60 | SWISSPEST19 | Swiss Pesticides and Metabolites ? 1,358
 - ▶ S61 | UJICCSLIB | Collision Cross Section (CCS) Library from UJI ? 574
 - ▶ S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag ? 258
 - ▶ S68 | HSDBTPS | Transformation Products Extracted from HSDB Content in PubChem ? 97

Transformation Products: Filling the Data Gaps!



PubChem NORMAN Suspect List Exchange

- ▼ NORMAN Suspect List Exchange Classification ? 113,080
- ▶ S13 | EUCOSMETICS | Combined Inventory of Ingredients Employed in Cosmetic Products (2000) and Revised Inventory (2006) ? 3,85
 - ▶ S25 | OECDPFAS | List of PFAS from the OECD ? 3,677
 - ▶ S36 | UBAPMT | Potential Persistent, Mobile and Toxic (PMT) substances ? 254
 - ▶ S50 | CCSCOMPEND | The Unified Collision Cross Section (CCS) Compendium ? 885
 - ▶ S60 | SWISSPEST19 | Swiss Pesticides and Metabolites from Kiefer et al 2019 ? 1,343
 - ▶ S61 | UJICCSLIB | Collision Cross Section (CCS) Library from UJI ? 574
 - ▶ S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag ? 258
 - ▶ S68 | HSDBTPS | Transformation Products Extracted from HSDB Content in PubChem ? 102
 - ▶ S69 | LUXPEST | Pesticide Screening List for Luxembourg ? 386
 - ▶ S72 | NTUPHTW | Pharmaceutically Active Substances from National Taiwan University ? 1,068
 - ▶ S75 | CyanoMetDB | Comprehensive database of secondary metabolites from cyanobacteria ? 2,088
 - S00 | SUSDAT | Merged NORMAN Suspect List: SusDat ? 99,130
 - S01 | MASSBANK | NORMAN Compounds in MassBank EU ? 7,164
 - S02 | STOFFIDENT | HSWT/LfU STOFF-IDENT Database of Water-Relevant Substances ? 11,261
 - S03 | NORMANCT15 | NORMAN Collaborative Trial Targets and Suspects ? 624
 - S04 | UJIBADE | Target List from UJI used in Bade et al 2015 ? 542

▼ Pharmacology and Biochemistry ? 112,039

▶ Human Metabolite Information ? 64,199

Metabolism/Metabolites ? 8,204

Transformations ? 5,857

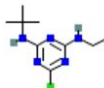
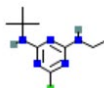
PubChem Terbutylazine (Compound)

8.5 Transformations

Page 3 of 25 items View More Rows & Details

Download

SORT BY Please Choose One

Predecessor Image	Predecessor Name	Transformation	Successor Image	Successor Name	Evidence DOI
	Terbutylazine	Mammalian metabolism		6-Chloro-1,3,5-triazine-2,4-diamine	10.5281/zenodo.382
	Terbutylazine	Deethylation		Terbutylazine-desethyl	10.1007/s13361-017-

Transformation Products: Filling the Data Gaps!

PubChem Terbutylazine (Compound)

7 Agrochemical Information

7.1 Agrochemical Category

Pesticides -> Herbicides -> [Triazine](#) herbicides -> Chlorotriazine herbicides

S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag | DOI:10.5281/zenodo.3754448

► [NORMAN Suspect List Exchange](#)

7.2 Agrochemical Transformations

Terbutylazine has known environmental transformation products that include [Terbutylazine-2-hydroxy](#), [Terbutylazine-desethyl](#), and [Terbutylazine-desethyl-2-hydroxy](#).

S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag | DOI:10.5281/zenodo.3754448

► [NORMAN Suspect List Exchange](#)

Terbutylazine has known environmental transformation products that include CSAA036479, CSAA04949, CSCD648241, CSCD692760, GS31398, MT1, GS 26379, MT13, GS 23158, Terbutylazine metabolite MT14, Terbutylazine metabolite MT23, and Terbutylazine metabolite MT24.

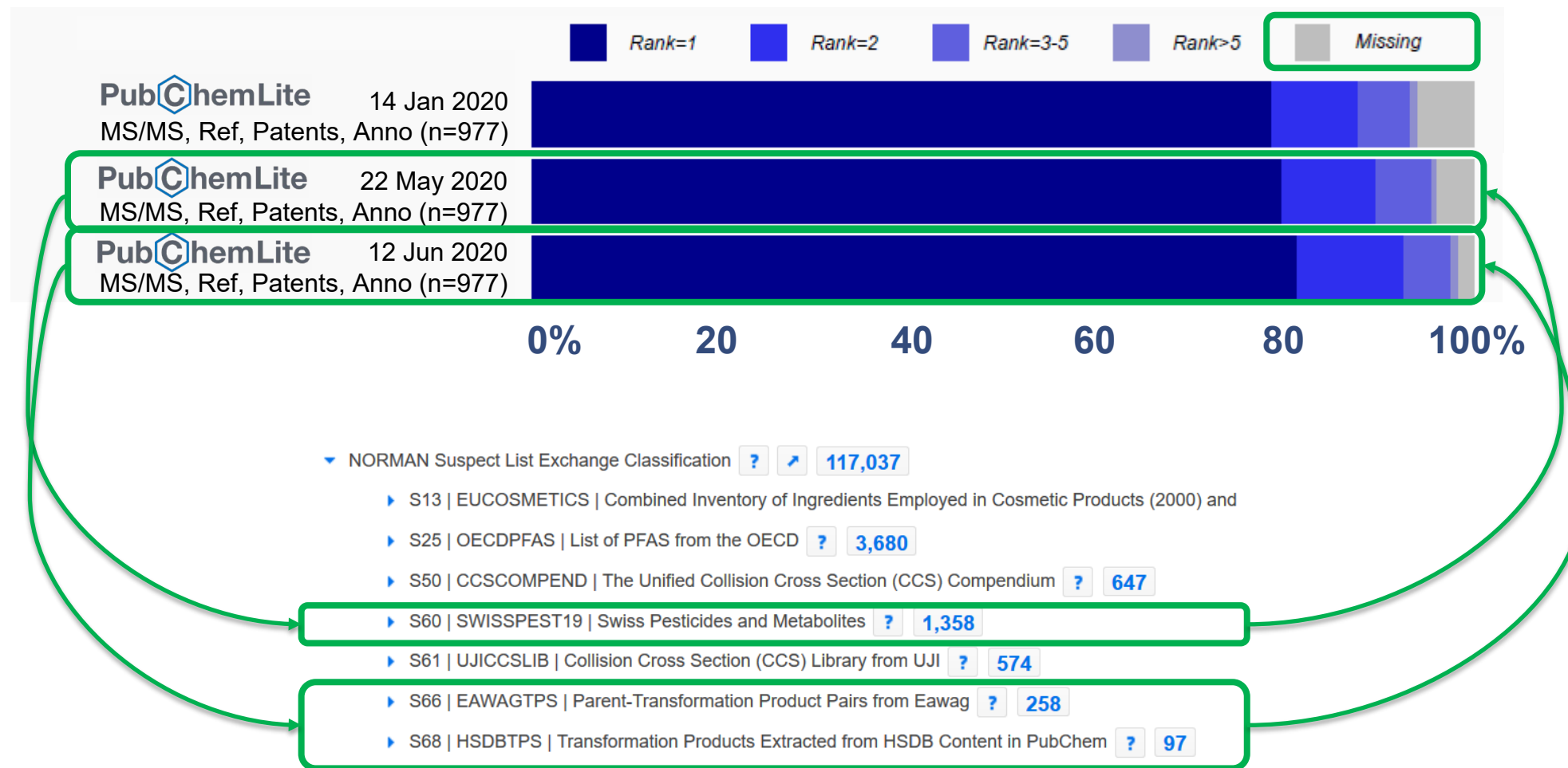
S60 | SWISSEST19 | Swiss Pesticides and Metabolites from Kiefer et al 2019 | DOI:10.5281/zenodo.3544759

► [NORMAN Suspect List Exchange](#)

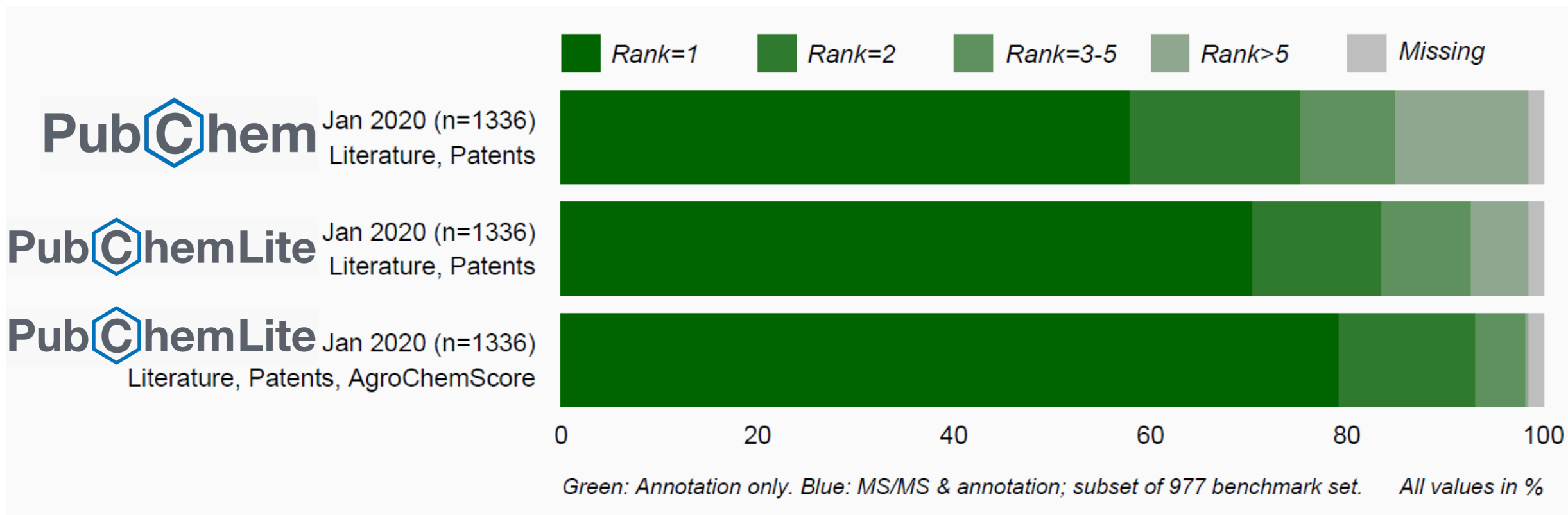
▼ Agrochemical Information	?	3,045
Agrochemical Category	?	1,903
Agrochemical Transformations	?	1,383
EU Pesticides Data	?	1,213
USDA Pesticide Data Program	?	579

PubChem Lite
EXPOSOMICS

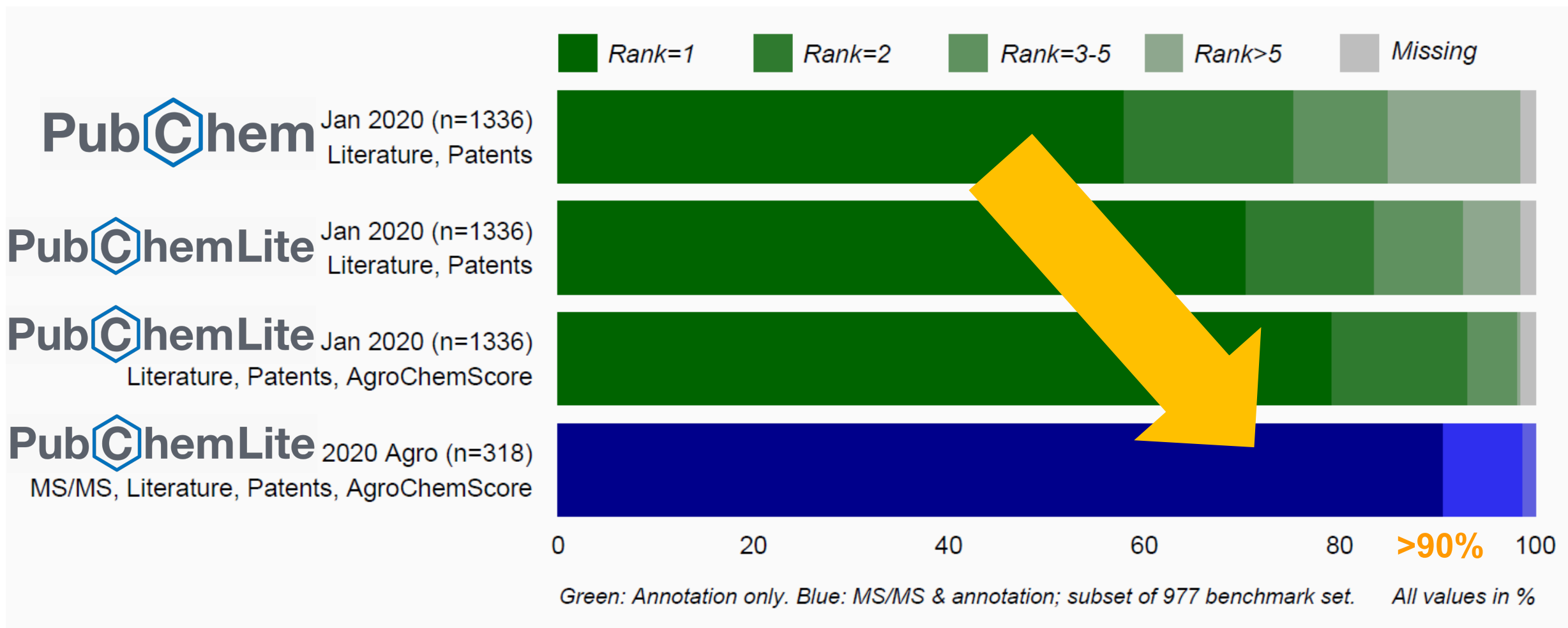
Assessing the Missing Entries in PubChemLite



Influence of the Annotation Content in PubChemLite



Influence of the Annotation Content in PubChemLite



Outline for today

- Background

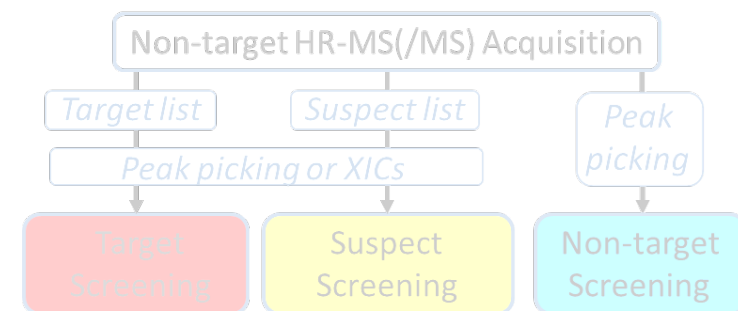
- Environmental cheminformatics & HR-MS
- Non-target screening and identification confidence
- Compound databases and spectral libraries

- Introduction to PubChemLite and MetFrag

- Examples (MassBank+PubChemLite+MetFrag)

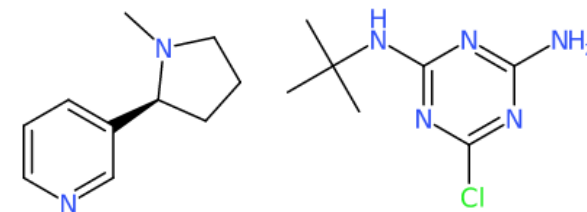
- Nicotine – relating identifications to disease outcomes
- Desethylterbutylazine – isobars and the role of metadata

- Closing – Perspectives and Community Contribution



PubChemLite
EXPOSOMICS

MetFrag



MetFrag Example: Nicotine and Disease Associations

<https://msbi.ipb-halle.de/MetFrag/>

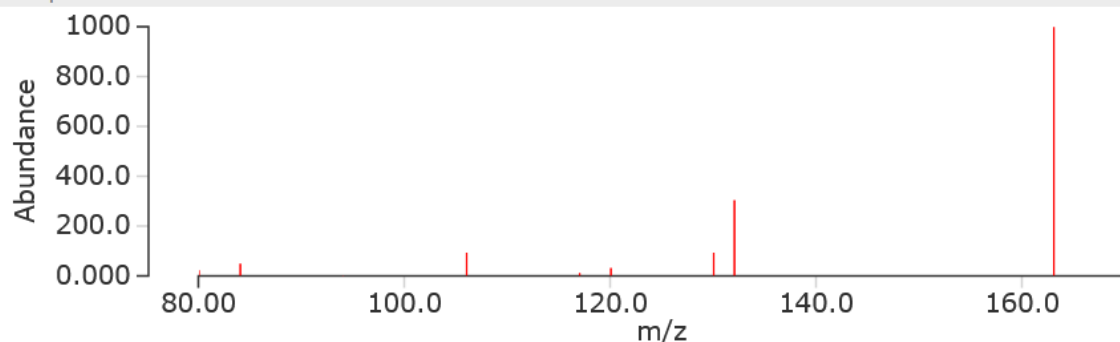
A screenshot of the MetFrag web interface. The header shows the "MetFrag" logo and the text "In silico fragmentation for computer assisted identification of metabolite mass spectra". Below this is a green bar labeled "Database Settings". The main form has fields for "Database:", "Neutral Mass:", "Formula:", and "Identifiers:". The "Database:" dropdown menu is open, showing a list of databases. A red arrow points from the "PubChemLite_01Jan2021_exposomics" option in the dropdown to the "PubChemLite EXPOSOMICS" logo in the top right corner. To the right of the database field is a "Parent Ion:" field with a dropdown menu set to "[M+H]+" and a "Calculate" button. Below the database field is a "Retrieve Candidate" button. At the bottom right is a "Download Candidates" button. At the bottom left is a "Candidate Filter & Score Settings" link.

MetFrag Example: Nicotine and Disease Associations

MassBank Record: EQ300801

Nicotine; LC-ESI-QFT; MS2; CE: 15; R=35000; [M+H]⁺

Mass Spectrum



[metabolomics-usi visualisation](#)

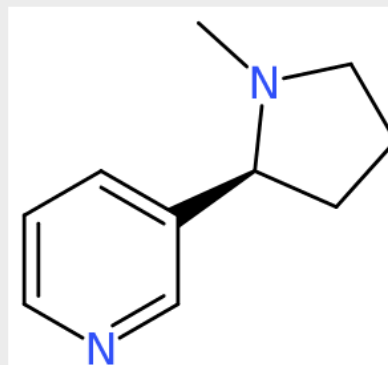
ACCESSION: EQ300801

RECORD_TITLE: Nicotine; LC-ESI-QFT; MS2; CE: 15; R=35000; [M+H]⁺

DATE: 2015.08.25

AUTHORS: Beck B, Stravs M, Schymanski E, Singer H, Department of Environmental

Chemical Structure



PK\$NUM_PEAK: 9

PK\$PEAK: m/z int. rel.int.

80.0494	6261028.7	23
84.0807	13197924.1	50
94.065	967625.9	3
106.065	24640249.3	93
117.0572	3192413.5	12
120.0807	8648923.7	32
130.0651	24669353.9	93
132.0807	80112590.7	304
163.1229	263120223.6	999

//



MetFrag Example: Nicotine and Disease Associations



MassBank
High Quality Mass Spectral Database

Database Settings

Database:

PubChemLite_01Jan2021

Parent Ion:

163.123

[M+H]⁺

Calculate

Neutral Mass:

162.11572

Search ppm:

5

Formula:

Identifiers:

Retrieve Candidates

Download Candidates

Candidate Filter & Score Settings

Fragmentation Settings & Processing

Mzppm:

5

Mzabs:

0.001

Mode:

[M+H]⁺

Tree depth:

2

Group candidates

☒

MS\$FOCUSSED_ION: BASE_PEAK 163.1229
MS\$FOCUSSED_ION: PRECURSOR_M/Z 163.123
MS\$FOCUSSED_ION: PRECURSOR_TYPE [M+H]⁺

MS/MS Peak list

```
80.0494 6261028.7 23
84.0807 13197924.1 50
94.065 967625.9 3
106.065 24640249.3 93
117.0572 3192413.5 12
120.0807 8648923.7 32
130.0651 24669353.9 93
132.0807 80112590.7
304
163.1229 263120223.6
999
```

Show Spectrum

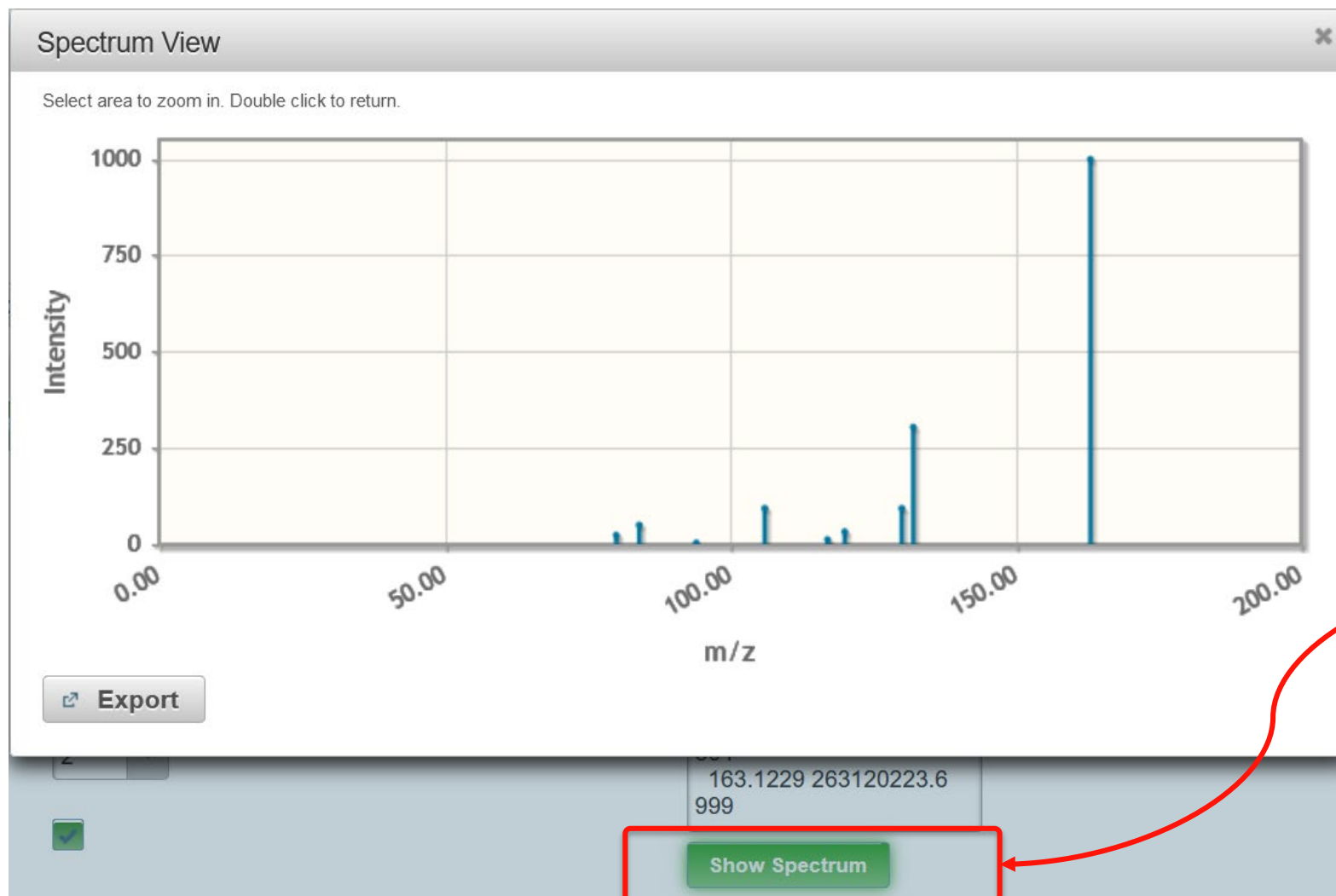
PK\$NUM_PEAK: 9
PK\$PEAK: m/z int. rel.int.
80.0494 6261028.7 23
84.0807 13197924.1 50
94.065 967625.9 3
106.065 24640249.3 93
117.0572 3192413.5 12
120.0807 8648923.7 32
130.0651 24669353.9 93
132.0807 80112590.7 304
163.1229 263120223.6 999
//



MetFrag Example: Nicotine and Disease Associations



Check the spectrum (to see it copied well)



PK\$NUM_PEAK: 9
PK\$PEAK: m/z int. rel.int.

80.0494	6261028.7	23
84.0807	13197924.1	50
94.065	967625.9	3
106.065	24640249.3	93
117.0572	3192413.5	12
120.0807	8648923.7	32
130.0651	24669353.9	93
132.0807	80112590.7	304
163.1229	263120223.6	999

//

MetFrag Example: Nicotine and Disease Associations



Database Settings

Database: PubChemLite_01Jan2021

Parent Ion: 163.123 [M+H]⁺ **Calculate**

Neutral Mass: 162.11572 Search ppm: 5

Formula:

Identifiers:

Retrieve Candidates 62 Candidates **Download Candidates**

Candidate Filter & Score Settings

Fragmentation Settings & Processing

Mzppm: 5

Mzabs: 0.001

Mode: [M+H]⁺

Tree depth: 2

Group candidates ☒

MS/MS Peak list

```
80.0494 6261028.7 23
84.0807 13197924.1 50
94.065 967625.9 3
106.065 24640249.3 93
117.0572 3192413.5 12
120.0807 8648923.7 32
130.0651 24669353.9 93
132.0807 80112590.7
304
163.1229 263120223.6
999
```

Show Spectrum



MetFrag Example: Nicotine and Disease Associations



MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS

Candidate Filter & Score Settings

Candidate Filters

- ☐ Element
- ☐ Element
- ☐ Substructure
- ☐ Substructure
- ☐ Substructure
- ☐ Substructure
- ☐ Minimum
- ☐ Maximum
- ☐ Suspect

MetFrag Scoring Terms

- ☐ Substructure Inclusion
- ☐ Substructure Exclusion
- ☐ Retention Time
- ☐ Suspect Inclusion Lists
- ☐ Spectral Similarity (MoNA)
- ☒ Exact Spectral Similarity (MoNA)
- ☐ Statistical Scoring

Database Scoring Terms

Select Item(s) 0 of 14 item(s) selected

Select Item(s) 4 of 14 item(s) selected

Fragmentation Settings & Processing



MetFrag Example: Nicotine and Disease Associations



Process the candidates!



Fragmentation Settings & Processing

Mzppm: 5

Mzabs: 0.001

Mode: [M+H]⁺

Tree depth: 2

Group candidates ☒

MS/MS Peak list

80.0494 6261028.7 23
84.0807 13197924.1 50
94.065 967625.9 3
106.065 24640249.3 93
117.0572 3192413.5 12
120.0807 8648923.7 32
130.0651 24669353.9 93
132.0807 80112590.7
304
163.1229 263120223.6
999

Show Spectrum

Download Parameters

Process Candidates

i 62 Candidates processed

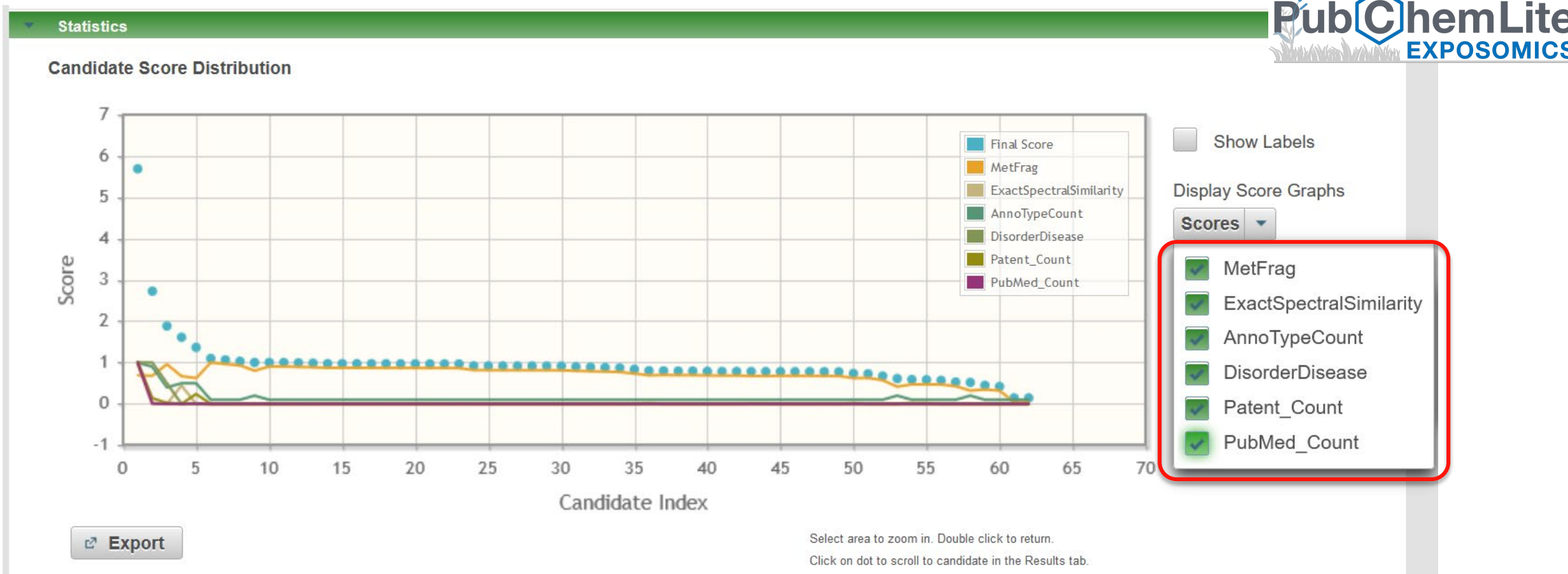
Statistics

Results

MetFrag Example: Nicotine and Disease Associations



Overview of all candidates: “Statistics” Plot



MetFrag Example: Nicotine and Disease Associations



MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS

Results

Weights

MetFrag (1st)	<input type="range"/>	100	%
ExactSpectralSimilarity (2nd)	<input type="range"/>	100	%
AnnoTypeCount (3rd)	<input type="range"/>	100	%
DisorderDisease (4th)	<input type="range"/>	100	%
Patent_Count (5th)	<input type="range"/>	100	%
PubMed_Count (6th)	<input type="range"/>	100	%

Experimental evidence / values

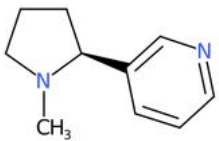
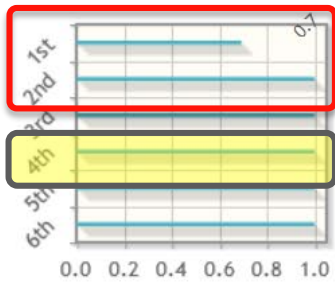
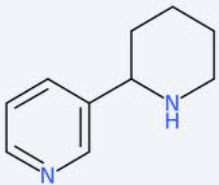
Download Results

Filter Candidates by explained MS/MS Peaks

MS/MS Peaks

Filter Candidates

Disease/Disorder information available

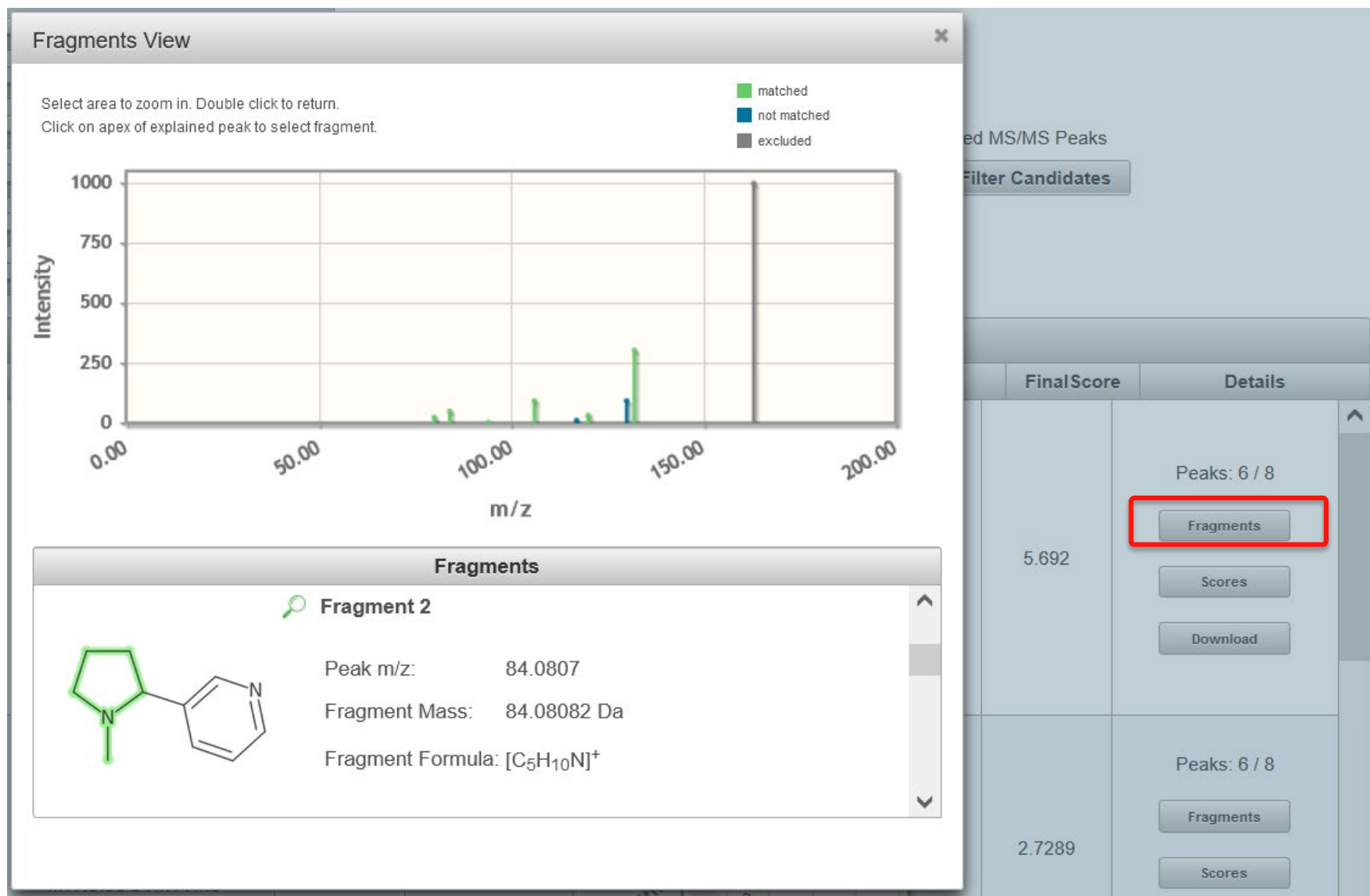
#	Molecule	Identifier	Mass	Formula	Normalized Scores	FinalScore	Details
1	 3-[(2S)-1-methylpyrrolidin-2-yl]pyridine	89594 InChIKeyBlock1 = SNICXCGAKADSCV	162.1157	C ₁₀ H ₁₄ N ₂		5.692	Peaks: 6 / 8 Fragments Scores Download
2	 3-piperidin-2-ylpyridine	2181 InChIKeyBlock1 = MTXSIJUGVMTTMU	162.1157	C ₁₀ H ₁₄ N ₂		2.7289	Peaks: 6 / 8 Fragments Scores Download

MetFrag Example: Nicotine and Disease Associations



MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS



MetFrag Example: Nicotine and Disease Associations



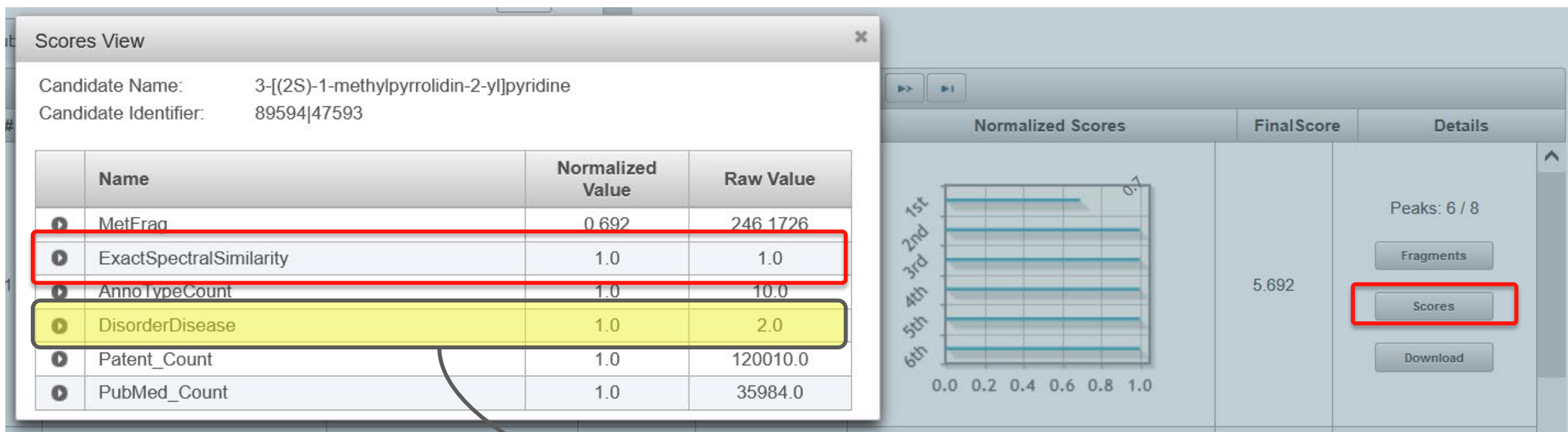
MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS

Level 2: Probable structure

- a) by library spectrum match
- b) by diagnostic evidence

MS, MS², Library MS² >0.9 sim
MS, MS², Exp. data



PubChem



MetFrag Example: Nicotine and Disease Associations



MassBank
High Quality Mass Spectral Database

PubChem
EXPOSOMICS

Results

Weights		
MetFrag (1st)	<input type="range"/>	100 %
ExactSpectralSimilarity (2nd)	<input type="range"/>	100 %
AnnoTypeCount (3rd)	<input type="range"/>	100 %
DisorderDisease (4th)	<input type="range"/>	100 %
Patent_Count (5th)	<input type="range"/>	100 %
PubMed_Count (6th)	<input type="range"/>	100 %

Download Results

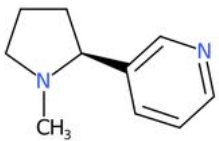
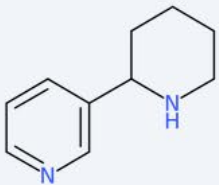
Filter Candidates by explained MS/MS Peaks

MS/MS Peaks

Filter Candidates

Disease/Disorder information available

1 2 3 4 5 6 7

#	Molecule	Identifier
1	 3-[(2S)-1-methylpyrrolidin-2-yl]pyridine	89594 InChIKeyBlock1 = SNICXCGAKADSCV
2	 3-piperidin-2-ylpyridine	2181 InChIKeyBlock1 = MTXSIJUGVMTTMU

PubChem About Blog Submit Contact

Explore Chemistry

Quickly find chemical information from authoritative sources

89594

6th
0.0 0.2 0.4 0.6 0.8 1.0

Download



PubChem Annotations: Disease Associations

PubChem Nicotine (Compound)

14 Associated Disorders and Diseases

246 items View More Rows & Details

Download

Cite

Download

Disease	Evidence Type	Evidence PMID
Abdominal Pain	marker/mechanism	9179092
Abnormalities, Drug-Induced	marker/mechanism	16193507
Acute Kidney Injury	marker/mechanism	21511693
Airway Obstruction	marker/mechanism	33082140
Akinetic Mutism	therapeutic	19455735
1 2 3 ... 50 Next >		

► [Comparative Toxicogenomics Database \(CTD\)](#)

Disease	References
	PubMed: 9137998 , 8294547 , 10636262 , 6890513 ,

CONTENTS

7 Drug and Medication Information

8 Agrochemical Information

9 Pharmacology and Biochemistry

10 Use and Manufacturing

11 Identification

12 Safety and Hazards

13 Toxicity

14 Associated Disorders and Diseases

Outline for today

- Background

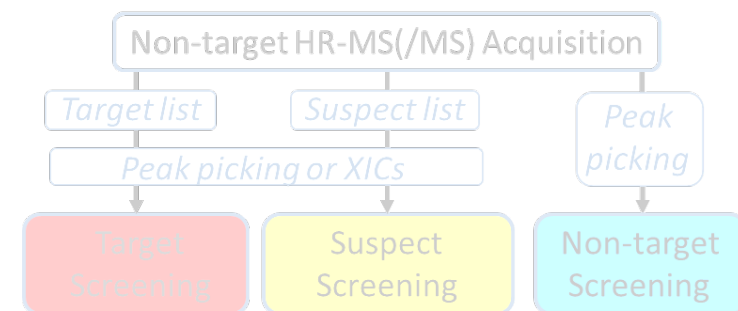
- Environmental cheminformatics & HR-MS
- Non-target screening and identification confidence
- Compound databases and spectral libraries

- Introduction to PubChemLite and MetFrag

- Examples (MassBank+PubChemLite+MetFrag)

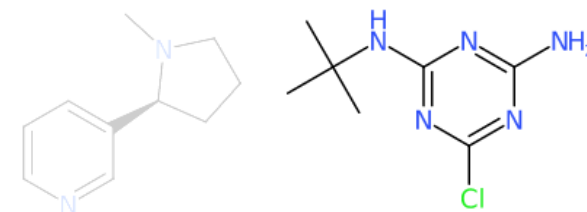
- Nicotine – relating identifications to disease outcomes
- Desethylterbutylazine – isobars and the role of metadata

- Closing – Perspectives and Community Contribution

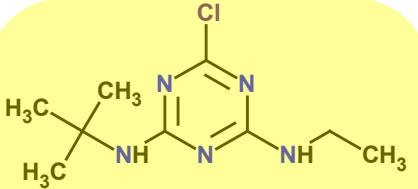
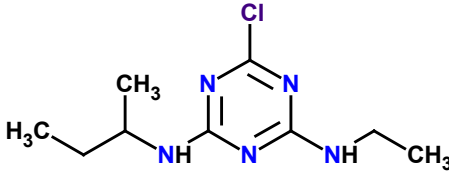
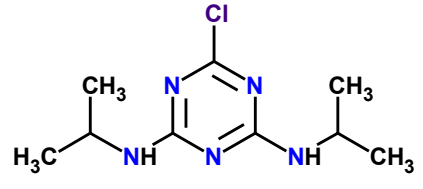
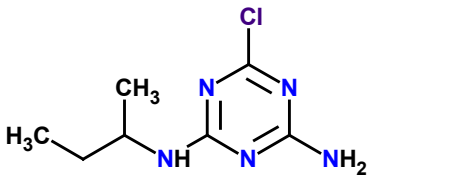
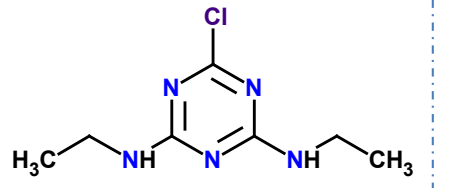
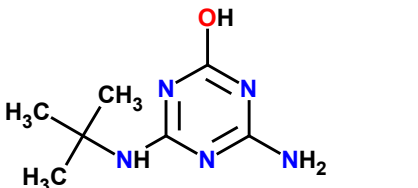
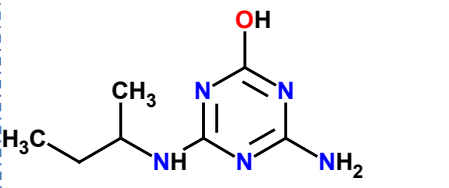
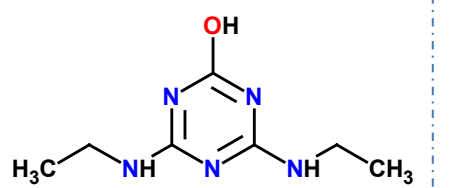


PubChemLite
EXPOSOMICS

MetFrag



MetFrag Example: Isobars and Metadata!

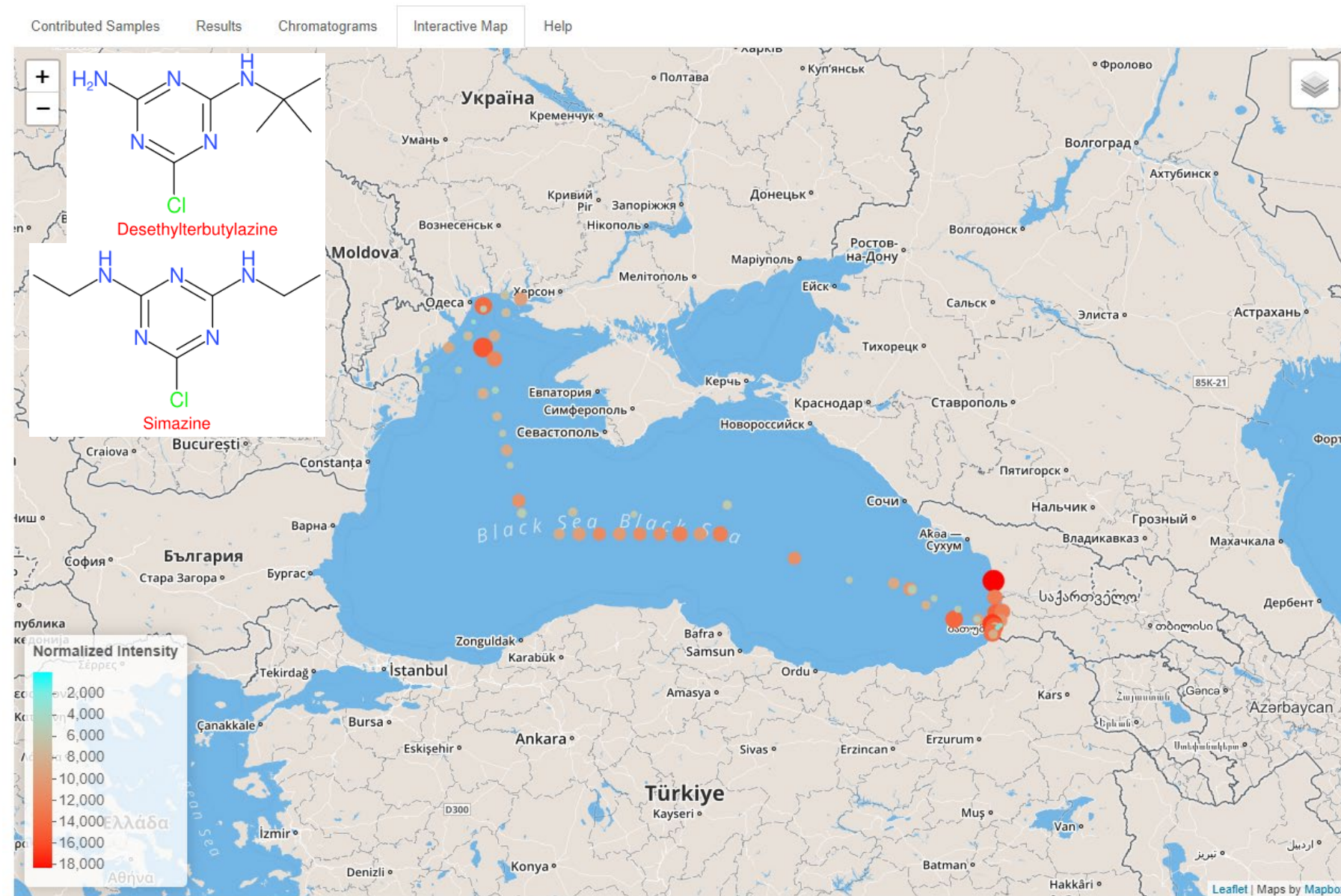
$C_9H_{16}ClN_5$ m/z 229.1094 Da	 Terbutylazine Detects: 12; # Refs: 220	 Sebutylazine Detects: 3; # Refs: 51	(no related compound at this mass)	 Propazine Detects: 3; # Refs: 201
$C_7H_{12}ClN_5$ m/z 201.0781 Da	 Terbutylazine-desethyl Detects: 9; # Refs: 92	 Sebutylazine-desethyl Detects: 1; # Refs: 14	 Simazine Detects: 4; # Refs: 518	(no related compound at this mass)
$C_7H_{13}N_5O$ m/z 183.1120 Da	 Terbutylazine-desethyl-2-hydroxy Detects: 2; # Refs: 57	 Sebutylazine-desethyl-2-hydroxy Detects: 0; # Refs: 3	 Simazine-2-hydroxy Detects: 2; # Refs: 66	(no related compound at this mass)

Schymanski *et al.* 2015, ABC,
 DOI: [10.1007/s00216-015-8681-7](https://doi.org/10.1007/s00216-015-8681-7)

MetFrag Example: Isobars and Metadata!

NORMAN Digital
Sample Freezing
Platform

Two very common
isomers ...



MetFrag Example: Isobars and Metadata!



MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS

Database Settings

Database: PubChemLite_01Jan2021

Parent Ion: 202.0854 [M+H]⁺ Calculate

Neutral Mass: 201.07812 Search ppm: 5

Formula:

Identifiers:

Retrieve Candidates 83 Candidates Download Candidates

Candidate Filter & Score Settings

Fragmentation Settings & Processing

Mzppm: 5

Mzabs: 0.001

Mode: [M+H]⁺

Tree depth: 2

Group candidates ☒

MS/MS Peak list

57.0699 34824.5 11
61.9792 27260 8
68.0244 499632.7 162
79.0059 755457.7 245
104.0011 904831.9 293
110.0462 250173.3 81
128.0568 46548.7 15
146.023 3077763.2 999

Show Spectrum

MassBank Record: EA067112



MetFrag Example: Isobars and Metadata!



MassBank
High Quality Mass Spectral Database

PubChemLite
EXPOSOMICS

Candidate Filter & Score Settings

Candidate Filters

- ☐ Element Inclusion
- ☐ Element Exclusion
- ☐ Substructure Inclusion
- ☐ Substructure Exclusion
- ☐ Substructure Information
- ☐ Minimum Number of Elements
- ☐ Maximum Number of Elements
- ☐ Suspect Inclusion Lists

Database Scoring Terms

Select Item(s) 3 of 14 item(s) selected

- ☐ AgroChemInfo
- ☒ AnnoTypeCount
- ☐ BioPathway
- ☐ DisorderDisease
- ☐ DrugMedicInfo
- ☐ FoodRelated
- ☐ Identification
- ☐ KnownUse
- ☒ Patent_Count
- ☐ PharmacolInfo
- ☒ PubMed_Count
- ☐ Related_CIDs
- ☐ SafetyInfo
- ☐ ToxicityInfo

MetFrag Scoring Terms

- ☐ Substructure Inclusion
- ☐ Substructure Exclusion
- ☐ Retention Time
- ☐ Suspect Inclusion Lists
- ☐ Spectral Similarity (MoNA)
- ☒ Exact Spectral Similarity (MoNA)
- ☐ Statistical Scoring

Database Scoring Terms

Select Item(s) 0 of 14 item(s) selected

Fragmentation Settings & Processing



MetFrag Example: Isobars and Metadata!



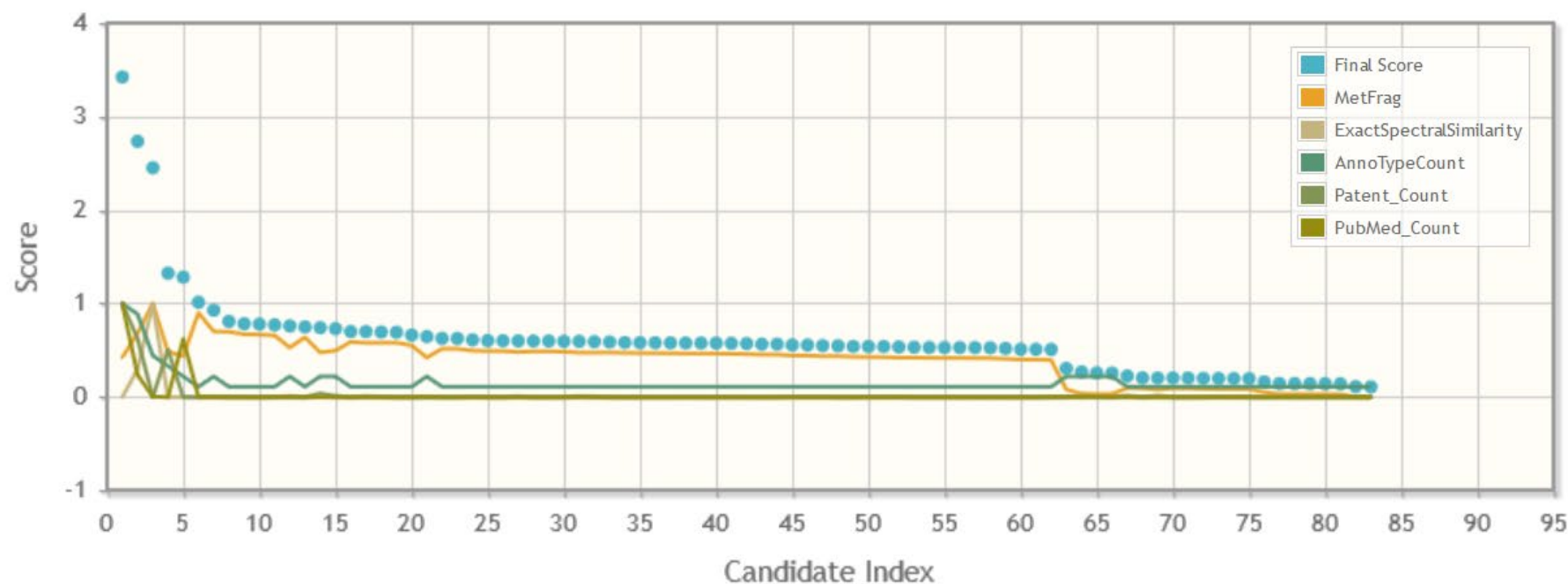
Process Candidates

83 Candidates processed

Download Parameters

Statistics

Candidate Score Distribution



Show Labels

Display Score Graphs

Scores

- ☒ MetFrag
- ☒ ExactSpectralSimilarity
- ☒ AnnoTypeCount
- ☒ Patent_Count
- ☒ PubMed_Count

Export

Select area to zoom in. Double click to return.

Click on dot to scroll to candidate in the Results tab.



MetFrag Example: Isobars and Metadata!



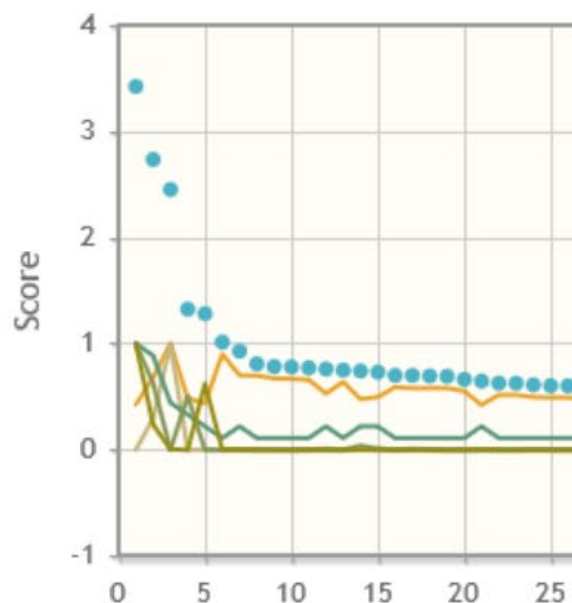
Process Candidates

83 Candidates processed

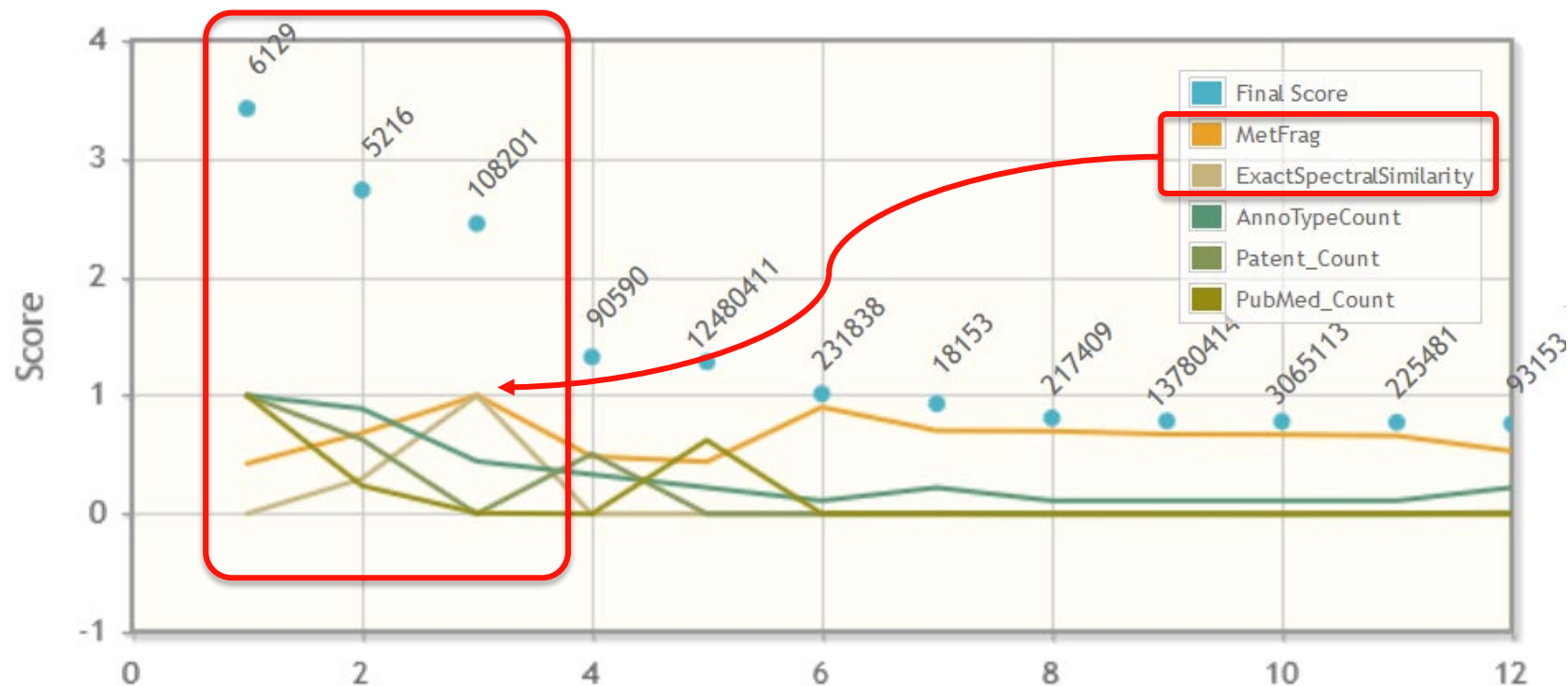
Download Parameters

Statistics

Candidate Score Distribution



Export



Click on dot to scroll to candidate in the Results tab.

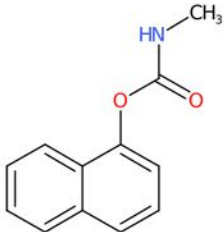
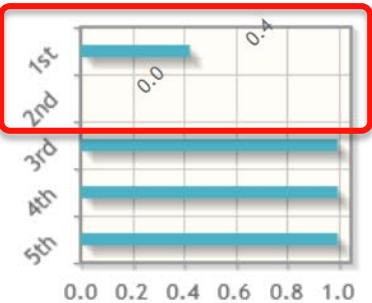
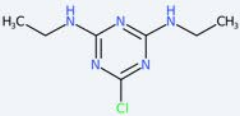
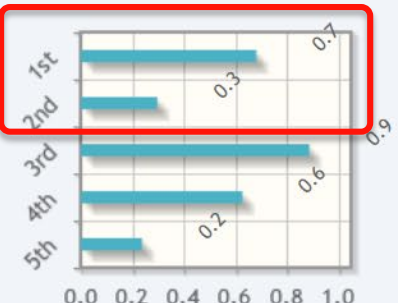
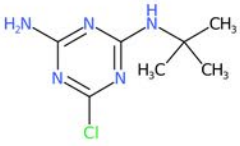
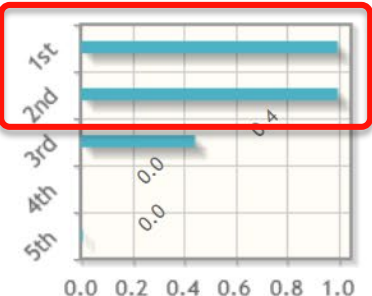


MetFrag Example: Isobars and Metadata!



MassBank
High Quality Mass Spectral Database

PubChem
EXPOSOMICS

#	Molecule	Identifier	Mass	Formula	Normalized Scores	FinalScore	Details
1	 naphthalen-1-yl N-methylcarbamate	6129 InChIKeyBlock1 = CVXBEEMKQHEXEN	201.07898	C ₁₂ H ₁₁ NO ₂		3.4241	Peaks: 2 / 8 Fragments Scores Download
2	 6-chloro-2-N,4-N-diethyl- 1,3,5-triazine-2,4- diamine	5216 InChIKeyBlock1 = ODCWYMIRDDJXKW	201.07812	C ₇ H ₁₂ ClN ₅		2.7357	Peaks: 4 / 8 Fragments Scores Download
3	 2-N-tert-butyl-6-chloro- 1,3,5-triazine-2,4-	108201 InChIKeyBlock1 = LMKQNTMFZLAJDV	201.07812	C ₇ H ₁₂ ClN ₅			Peaks: 6 / 8 Level 2: Probable structure a) by library spectrum match b) by diagnostic evidence

Spectral Match = 0, Frag 0.4

Spectral Match = 0.2976

Spectral Match = 1.000

Level 2: Probable structure
a) by library spectrum match
b) by diagnostic evidence

MetFrag Example: Isobars & Adjusted Weighting



MassBank
High Quality Mass Spectral Database

PubChem
EXPOSOMICS

Results

Weights

MetFrag (1st)	<input type="range"/>	100	%
ExactSpectralSimilarity (2nd)	<input type="range"/>	100	%
AnnoTypeCount (3rd)	<input type="range"/>	0	%
Patent_Count (4th)	<input type="range"/>	0	%
PubMed_Count (5th)	<input type="range"/>	0	%

Download Results

Filter Candidates by explained MS/MS Peaks

MS/MS Peaks **Filter Candidates**

#	Molecule	Identifier	Mass	Formula	Normalized Scores	FinalScore	Details
1	 2-N-tert-butyl-6-chloro-1,3,5-triazine-2,4-diamine	108201 InChIKeyBlock1 = LMKQNTMFZLAJDV	201.07812	C ₇ H ₁₂ ClN ₅		2.0	Level 2: Probable structure a) by library spectrum match b) by diagnostic evidence
2	 6-chloro-2-N,4-N-diethyl-1,3,5-triazine-2,4-diamine	5216 InChIKeyBlock1 = ODCWYMIRDDJXKW	201.07812	C ₇ H ₁₂ ClN ₅		0.98	Peaks: 4 / 8 Fragments Scores Download

Spectral Match = 1.000

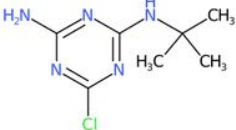
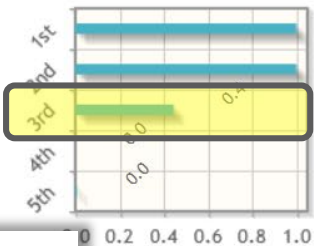
Spectral Match = 0.2976

MetFrag & Isobars: Supporting Annotation Content



MassBank
High Quality Mass Spectral Database

PubChem
EXPOSOMICS

#	Molecule	Identifier	Mass	Formula	Normalized Scores	FinalScore	Details
1		108201 InChIKey: Block1 = LMKQNTMEZLAJDV	201.07812	C ₇ H ₁₂ ClN ₅		2.0	Peaks: 6 / 8 Fragments Level 2: Probable structure a) by library spectrum match b) by diagnostic evidence

PubChem Desethylterbuthylazine (Compound)

7 Agrochemical Information

7.1 Agrochemical Category

Pesticides -> Herbicides -> Triazine herbicides -> Chlorotriazine herbicides -> Transformation products

S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag | DOI:10.5281/zenodo.3754448

► NORMAN Suspect List Exchange

7.2 Agrochemical Transformations

Terbutylazine-desethyl has known environmental transformation products that include Terbutylazine-desethyl-2-hydroxy.

Terbutylazine-desethyl is a known environmental transformation product of Terbutylazine.

S66 | EAWAGTPS | Parent-Transformation Product Pairs from Eawag | DOI:10.5281/zenodo.3754448

► NORMAN Suspect List Exchange

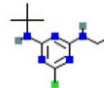
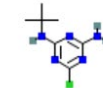
PubChem Desethylterbuthylazine (Compound)

8.1 Transformations

7 items View More Rows & Details

Download

SORT BY Please Choose One

Predecessor	Predecessor Name	Successor	Successor Name	Transformation	Enzyme
	terbuthylazine		desethyl-terbuthylazine	Environmental	



Outline for today

- Background

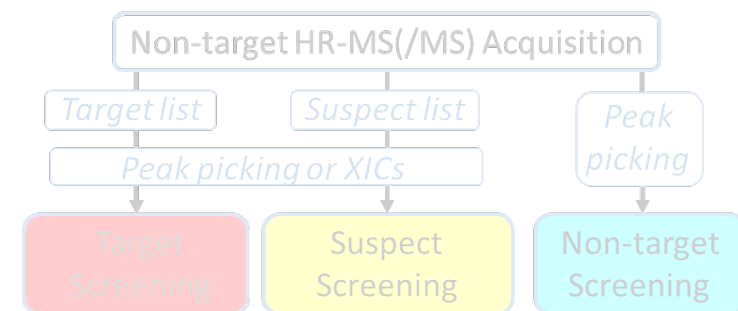
- Environmental cheminformatics & HR-MS
- Non-target screening and identification confidence
- Compound databases and spectral libraries

- Introduction to PubChemLite and MetFrag

- Examples (MassBank+PubChemLite+MetFrag)

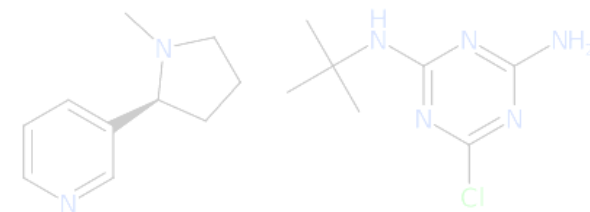
- Nicotine – relating identifications to disease outcomes
- Desethylterbutylazine – isobars and the role of metadata

- Closing – Perspectives and Community Contribution



PubChemLite
EXPOSOMICS

MetFrag



Expert Knowledge is YOUR Knowledge!

<https://ftp.ncbi.nlm.nih.gov/pubchem/Other/Submissions/>

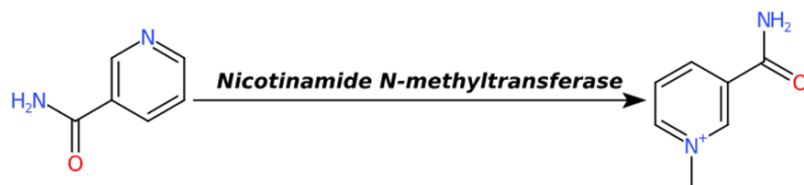
- Help us help you! Add data with FAIR templates

- Chemical Structures

PubChem_CID	Name	SMILES	InChIKey
2256	Atrazine	CCNC1=NC(=NC(=N1)Cl)NC(C)C	MXWJVTOOROXGIU-UHFFFAOYSA-N
2328	Bentazone	CC(C)N1C(=O)C2=CC=CC=C2NS1(=O)=O	ZOMSMJKLGFBRBS-UHFFFAOYSA-N
3030	Dicamba	COC1=C(C=CC(=C1C(=O)O)Cl)Cl	IWEDIXLBFLAXBO-UHFFFAOYSA-N
3120	Diuron	CN(C)C(=O)NC1=CC(=C(C=C1)Cl)Cl	XMTQQYYKAHVGBJ-UHFFFAOYSA-N

- Transformations

Predecessor_CID	Predecessor_Name	Transformation	Successor_CID	Successor_Name	Biosystem
13101	6PPD	Ozone	154926030	6PPD-quinone	Environment
2256	Atrazine	Environmental	13878	Deisopropyl-atrazine	Soil
2256	Atrazine	Mammalian metabolism	135408770	Ammeline	Mammal
2256	Atrazine	Fungal metabolism	22563	Desethyl-atrazine	Fungus



Nicotinamide to MNAM DOI:10.1124/dmd.112.049734

FAIR chemical structures in the Journal of Cheminformatics

Emma L. Schymanski and Evan E. Bolton

Letter to the Editor | 7 July 2021

i The [Letter Response to this article](#) has been published in *Journal of Cheminformatics* 2021 **13**:49

Reply to "FAIR chemical structure in the Journal of Cheminformatics"

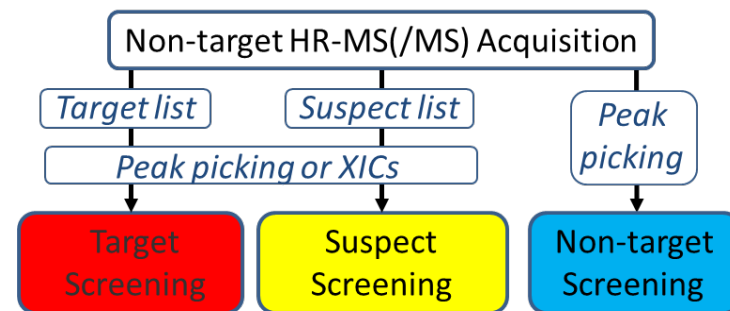
Rajarshi Guha, Nina Jeliaskova, Egon Willighagen and Barbara Zdrazil

Letter Response | 7 July 2021

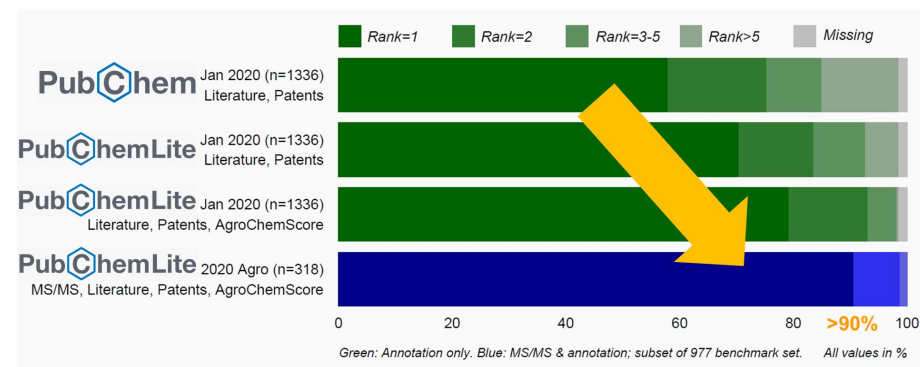
i The [Letter to the Editor to this article](#) has been published in *Journal of Cheminformatics* 2021 **13**:50

“Take home” Messages

- Background of HR-MS and Cheminformatics
- Introduction to MetFrag and PubChemLite
 - Mass spectral libraries help deliver Level 2a IDs
 - Annotation content is extremely powerful
- Please try the two examples & interpretation
 - More examples in the additional exercises
- Consider contributing your knowledge!



 **MassBank**
High Quality Mass Spectral Database



Thank you!

PubChemLite EXPOSOMICS

Email: emma.schymanski@uni.lu

Twitter: [@Eschymanski](https://twitter.com/Eschymanski)

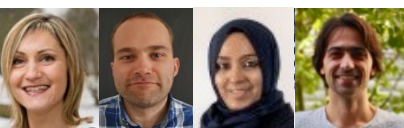
Slides @ [10.5281/zenodo.5115498](https://zenodo.org/record/5115498)

<https://msbi.ipb-halle.de/MetFrag/>

<https://pubchem.ncbi.nlm.nih.gov/>

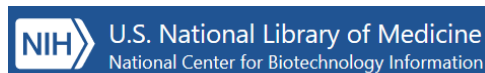
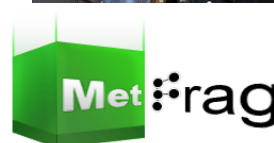
<https://massbank.eu/MassBank/>

https://www.wen.uni.lu/lcsb/research/environmental_cheminformatics/



PubChem

... and
team



Luxembourg National
Research Fund

