



PMI SCIENCE
PHILIP MORRIS INTERNATIONAL

High-Resolution Mass Spectrometry in Combination with *In Silico* Prediction Tools to Improve Accuracy for Compound Identification

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Philip Morris International R&D



Reduced-Risk Products (“RRPs”) is the term we use to refer to products that present, are likely to present, or have the potential to present less risk of harm to smokers who switch to these products versus continued smoking.

We have a range of RRP's in various stages of development, scientific assessment and commercialization. Because our RRP's do not burn tobacco, they produce far lower quantities of harmful and potentially harmful compounds than found in cigarette smoke.

Outline

- Challenge for complex matrix characterization
- GC-HR-MS instrumentation
- Linear retention index (LRI) modeling
- Compound identification in complex matrices
- *In silico* prediction (use of fragmentation software)
- Improvement in compound identification
- Conclusion

Chemical Characterization of Diverse Samples is Challenging

- More than 6'000 chemicals reported as present in tobacco plant and smoke ¹
- Many possible flavor compounds used in e-liquids or smoking products ^{2,3}



¹ The Chemical Components of Tobacco and Tobacco Smoke, A. Rodgman, T.A. Perfetti, 2013, 2nd Ed. CRC press.

² Leffingwell, J. C.; Young, H. J.; Bernasek, E. Tobacco flavoring for Smoking Products, R. J. Reynolds Tobacco Company, Winston-Salem, 1972.

³ EFSA flavoring substances database

Analytical Technique: GC-High Resolution (GC-HR-MS)

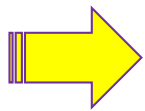
GC-HR-MS_1
(7200A Agilent Q-TOF-MS)

Volatiles and semi-
volatiles
LRI from 500 to 1'900



GC-HR-MS_2
(7200B Agilent Q-TOF-MS)

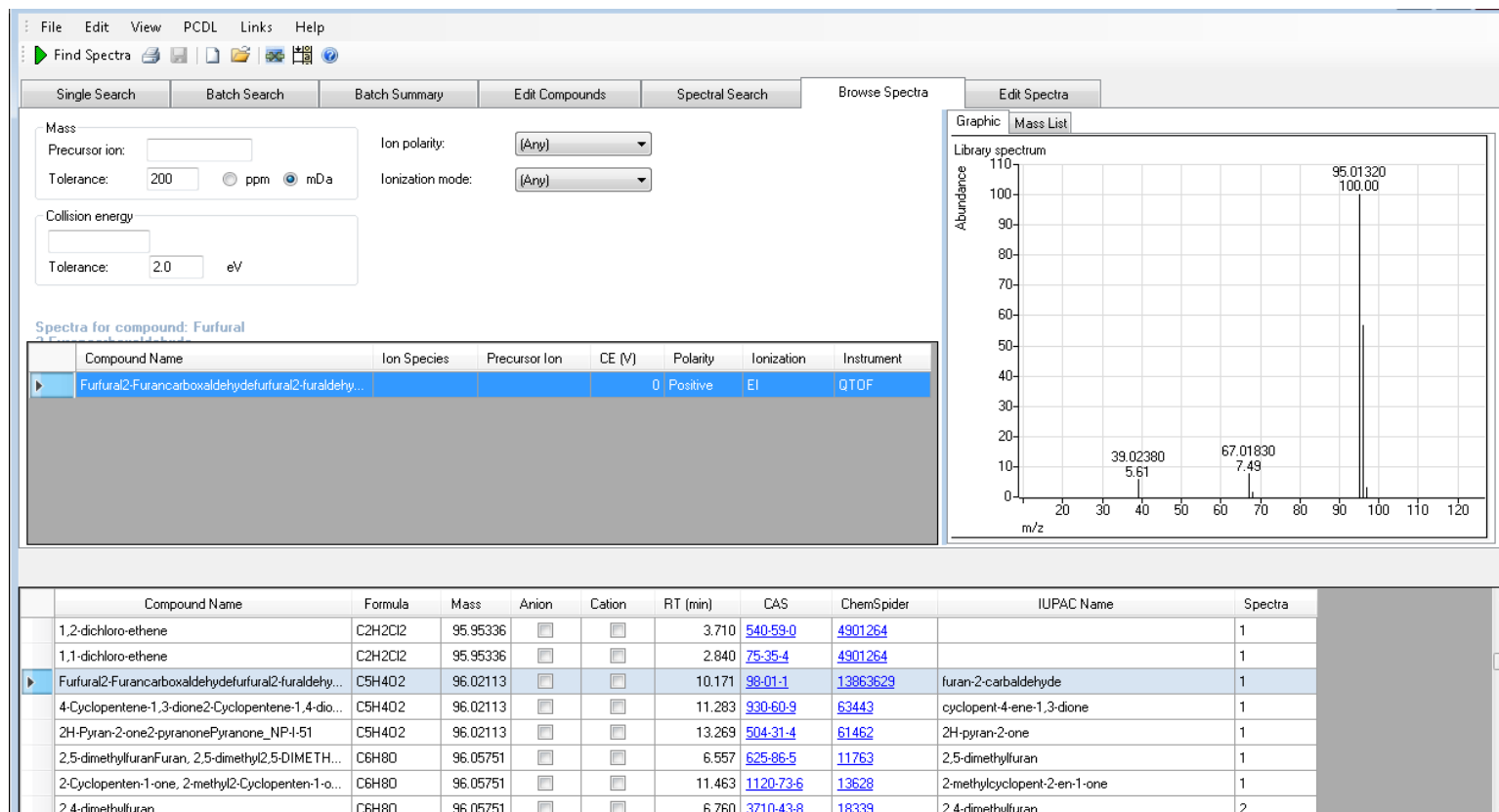
Apolar and polar
LRI from 1'000 to 3'000



Goal is to screen the broadest range of smoke constituents
Non-Targeted Screening

Volatile, Semi-volatile and Polar Compound Library

- 822 reference compounds were analyzed by GC-HR-MS
- Accurate mass spectra associated with their Linear Retention Indices were registered in the Personal Compound Database Library



An accurate LRI calculation and high quality spectra is key!!!

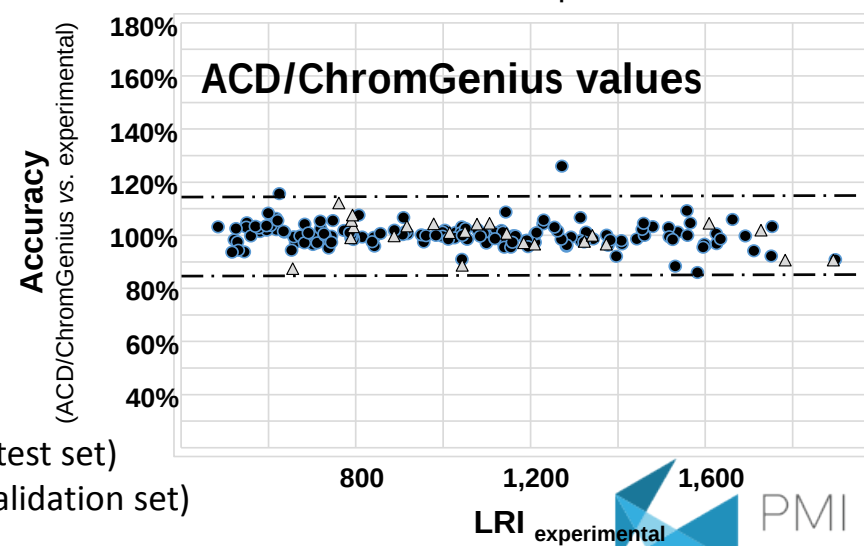
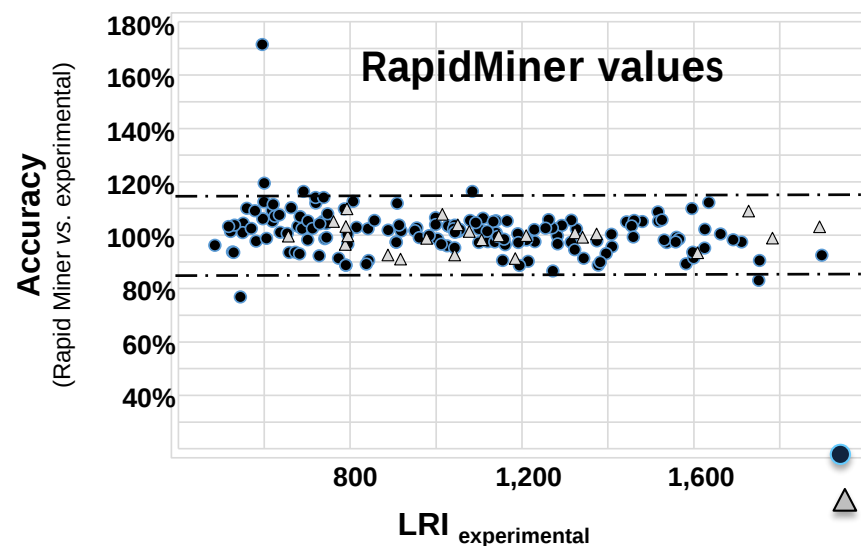
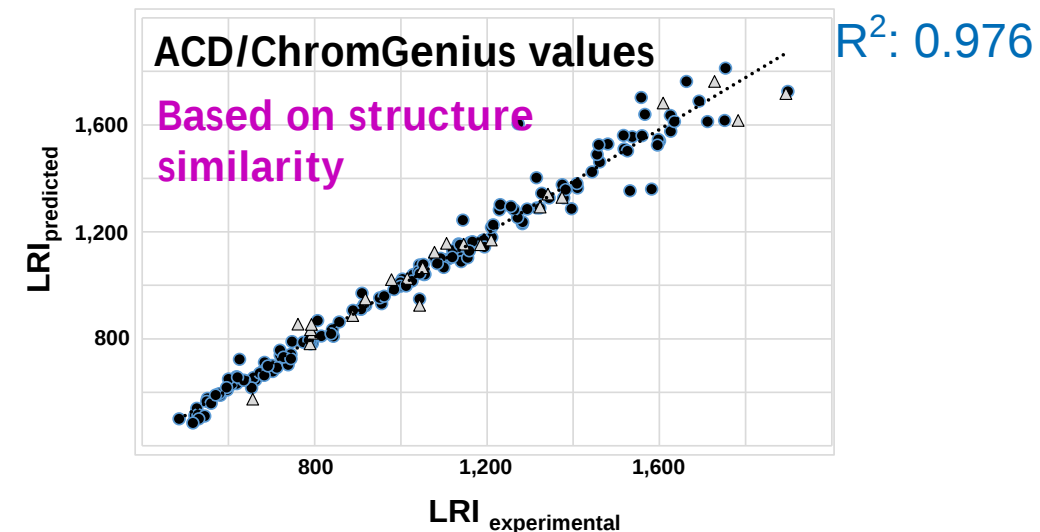
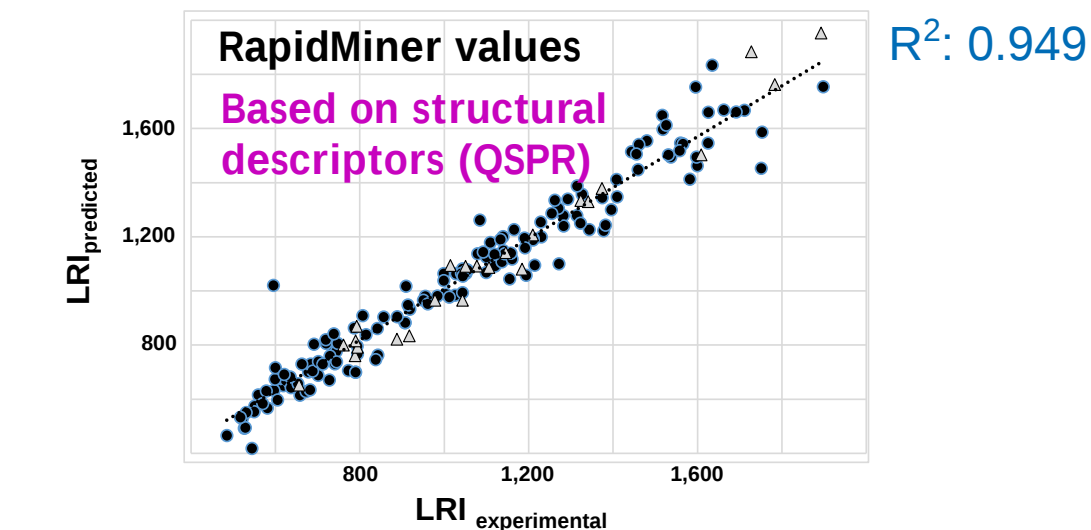
LRI Model Creation

- To predict LRI, two software combination were used:
 1. RapidMiner-Dragon
 2. ACD/Labs ChromGenius
- To build a relevant LRI prediction system 552 molecules were used:
 1. Experimental LRI
 2. Quantitative Structure-Property Relationship (QSPR) and structure similarities
- The experimental linear retention indices were randomly split as training (n=401) and test (n=151) sets
- Validation set (n=23) confirmed the great performance of both prediction models
- Discrepancy between two LRI predictions can highlight possible errors in the predicted values





Assessment of the Prediction Models (Test & Validation Sets)



- n=151 reference standards (test set)
- △ n=23 reference standards (validation set)

E. Dossin *et al.*, *Anal. Chem.* **2016**, *88*, 15, p7539-7547.

Unique Compound & Spectra Database (UCSD)

The screenshot displays the UCSD web interface for the compound PMI0009992. The top navigation bar includes 'Search Molecules', 'Submit New Molecule', and 'Administration'. The breadcrumb trail shows 'Search Molecules > Molecules > Molecule Details'. The main content area is divided into several panels:

- Chemical classification:** Thiophene, 1 - 1
- Generated Names / Structure Codes:**
 - Smiles: Cc1ccsc1C
 - InChI: InChI=1S/C6H8S/c1-5-3-7-4-6(5)/2/h3-4H,1-2H3
 - IUPAC: 3,4-dimethylthiophene
- Physical Measurements:** Ref. OR, Ref. BP, Ref. MP
- Calculated Mol. Properties:**
 - Mol. Formula: C6 H8 S
 - Mol. Weight: 112.19272 g/mol
 - Isotopic Mass: 112.03487 g/mol
 - ACD/LogP: 2.8
 - Solubility (w): 0.01 mol/L
 - H. Bond Acceptors: 0
 - H. Bond Donor: 0
 - Drug-like: Yes
 - Lead-like: Yes
- Predicted ADMET Properties:**
 - Hum. Intestinal Abs.: Moderate
 - Blood Brain Bar. Penetr.: High
 - Plasma Protein Binding: Non binder
 - CYP2D6 Inhibitor: No
 - Hepatotoxicity: Toxic
- Mass Spectra:** A plot showing relative intensity versus m/z.
- Alternatif Identifiers:**
 - Common Name: 3,4-Dimethylthiophene, 3,4-dimethylthiophene, Thiophene, 3,4-dimethyl-
 - CAS: 632-15-5, 1 - 1

On the left, a 'Batch List' section explains the hierarchy: molecule (neutral compound, e.g. PMI0000001), substance (molecule + counter ion, hydrate e.g. PMI0000001-A), and batches (physical or literature substances, e.g. BC0000014). It shows a tree structure for PMI0009992, including sub-batches like BC000014260 and BC000014291 (SPR000002647).

UCSD is our in-house database that contains:

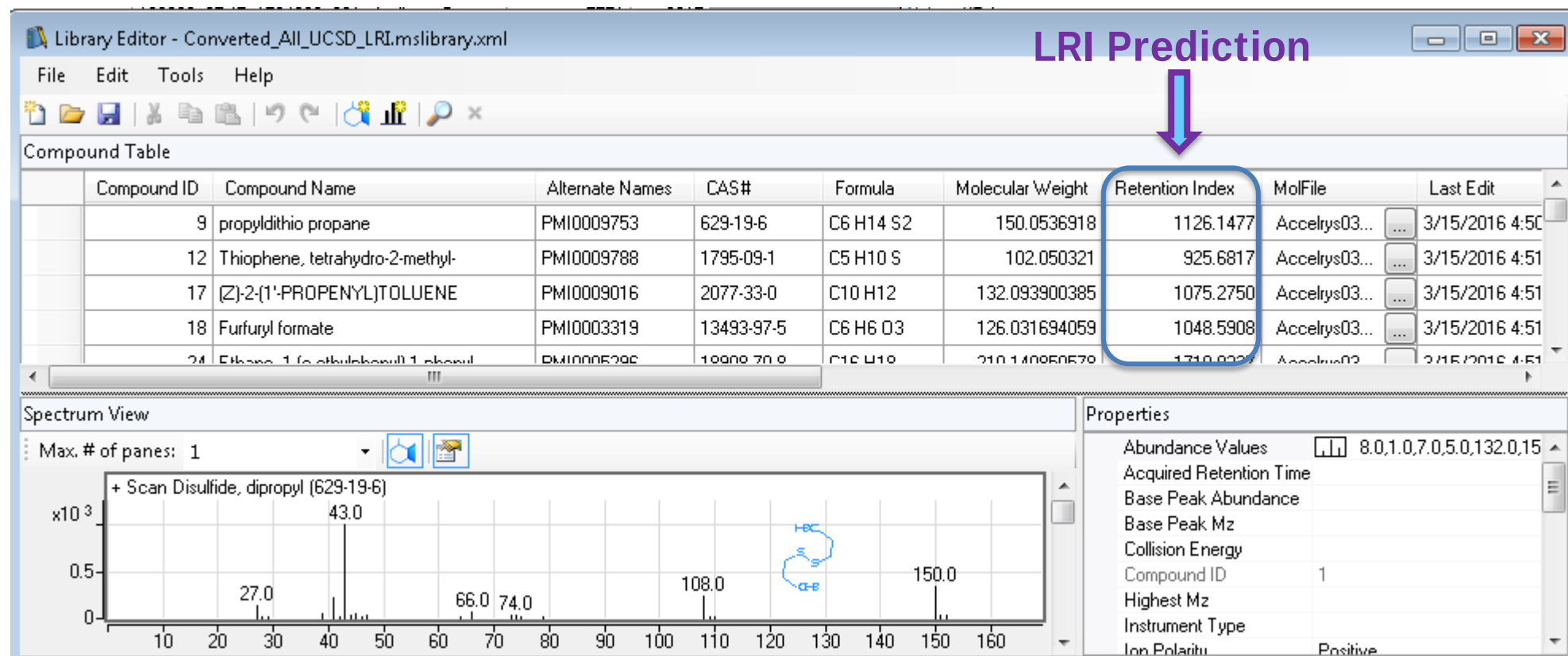
- 11'567 molecules
- 1'013 accurate mass spectra and LRI_{exp}

The 2 models were used to predict the LRI values of UCSD database (suspect list)

Our GC-MS conditions are suitable to analyze potentially 6'053 molecules

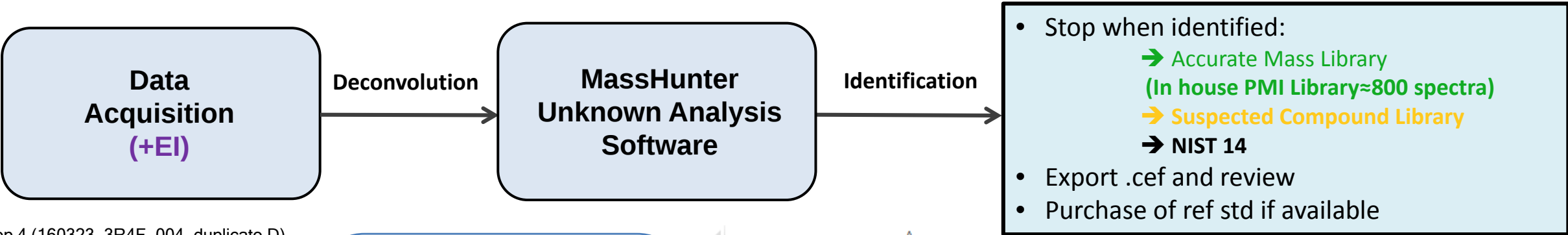
Of these 6'053 molecules, **3'646** have an available nominal EI Mass Spectra (NIST or Wiley)

Creation of the Suspected Compound Library

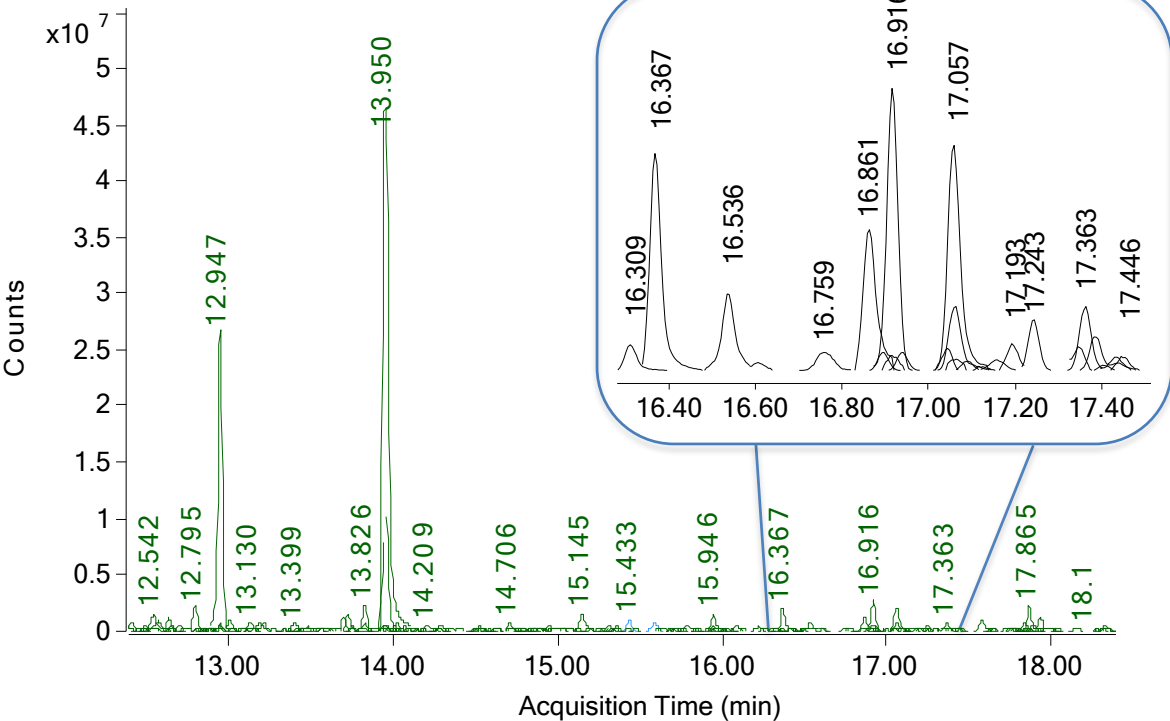


LRI prediction values were associated with the 3'646 nominal EI mass spectra extracted from commercial libraries

Compound Identification in Complex Matrices

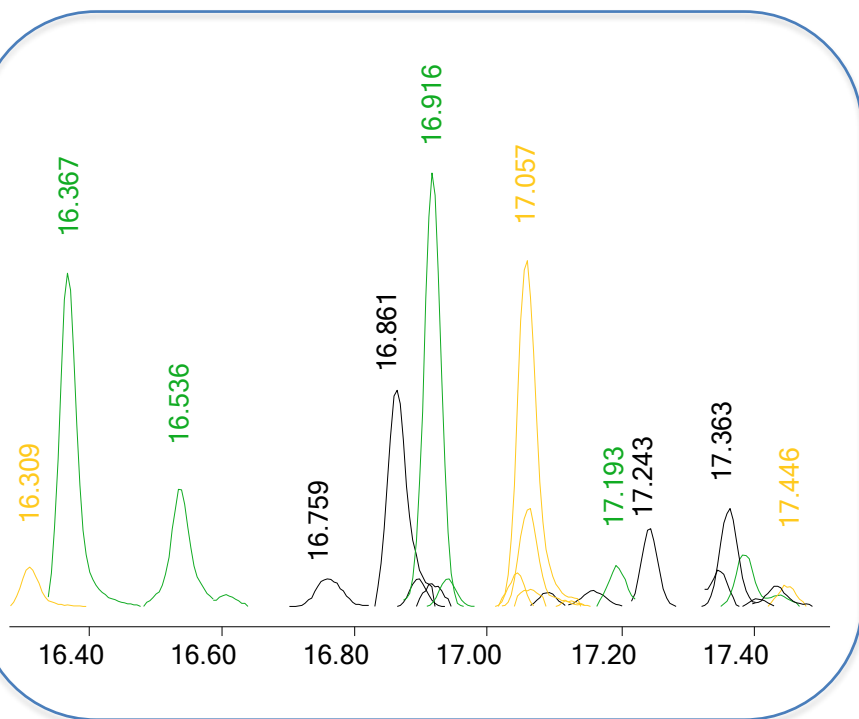


3R4F_Rep 4 (160323_3R4F_004_duplicate.D)



	A	B	C	D	E	F
1	Compound Name	Formula	CAS#	RT	Match Factor	Library File
378	2,5-Pyrrolidinedione (CAS)	C4H5NO2	123-56-8	16.309	67.5	UCSD
379	3-Hydroxypyridine	C5H5NO	109-00-2	16.367	52.8	FingerprintingDB624
380	4-Ethylphenol	C8H10O	123-07-9	16.536	76.4	FingerprintingDB624
381	3-Ethyl phenol	C8H10O	620-17-7	16.605	66.3	FingerprintingDB624
382	4-Fluorobenzoic acid, hex-4-yn-3-yl ester	C13H13FO2	1000299-15-3	16.759	61.1	NIST14
383	Butanoic acid, 3-methyl-, 3-methyl-3-butenyl ester	C10H18O2	54410-94-5	16.862	65.5	NIST14
384	Phenol, 3-amino-	C6H7NO	591-27-5	16.894	60.7	NIST14
385	Sarcosylsarcosine, N-methoxycarbonyl-, heptyl ester	C15H28N2O5	1000314-83-0	16.914	66.4	NIST14
386	Tridecane	C13H28	629-50-5	16.916	74.0	FingerprintingDB624
387	3,4-Dimethylphenol	C8H10O	95-65-8	16.934	60.3	FingerprintingDB624
388	4-Methyl-1,2-dihydronaphthalene	C11H12	4373-13-1	17.044	76.8	UCSD
389	Pentanedioic acid, 2-oxo-, dimethyl ester (CAS)	C7H10O5	13192-04-6	17.057	49.4	UCSD
390	Nonanoic acid (CAS)	C9H18O2	112-05-0	17.061	54.4	UCSD
391	Cyclohexasiloxane, dodecamethyl-	C12H36O6Si6	540-97-6	17.158	60.7	NIST14
392	Phenol, 2,3,4-trimethyl	C9H12O	526-85-2	17.193	46.4	FingerprintingDB624
393	Succinic acid, 2,5-dimethylphenyl ethyl ester	C14H18O4	1000329-86-0	17.243	56.4	NIST14
394	Naphthalene, 1,2-dihydro-3-methyl- (CAS)	C11H12	2717-44-4	17.347	65.5	UCSD
395	1,4:3,6-Dianhydro-.alpha.-d-glucopyranose	C6H8O4	1000098-14-8	17.363	77.8	NIST14
396	4-isopropylphenol	C9H12O	99-89-8	17.385	47.4	FingerprintingDB624
397	Divinylmethyl(chloromethyl)silane	C6H11ClSi	25202-02-2	17.433	56.1	NIST14
398	1H-Pyrrole-2,5-dione, 3-ethyl-4-methyl- (CAS)	C7H9NO2	20189-42-8	17.446	53.3	UCSD

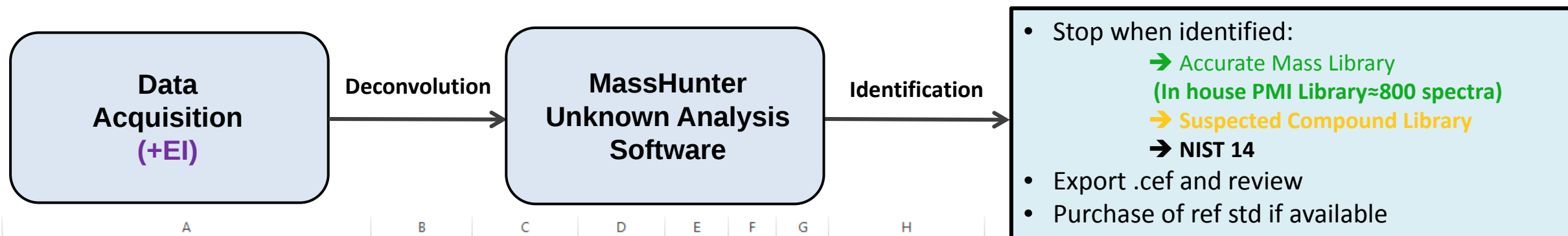
Reviewing Step



1	Compound Name	Formula	CAS#	Component RI	Library RI	Δ RI	Match Factor	Library File
378	2,5-Pyrrolidinedione (CAS)	C4H5NO2	123-56-8	1260.70	1249.71	-10.99	67.5	UCSD
379	3-Hydroxypyridine	C5H5NO	109-00-2	1264.45	1272.07	7.62	52.8	FingerprintingDB624
380	4-Ethylphenol	C8H10O	123-07-9	1275.38	1279.99	4.61	76.4	FingerprintingDB624
381	3-Ethyl phenol	C8H10O	620-17-7	1279.84	1282.00	2.16	66.3	FingerprintingDB624
382	4-Fluorobenzoic acid, hex-4-yn-3-yl ester	C13H13FO2	1000299-15-3	1289.82			61.1	NIST14
383	Butanoic acid, 3-methyl-, 3-methyl-3-butenyl ester	C10H18O2	54410-94-5	1296.49			65.5	NIST14
384	Phenol, 3-amino-	C6H7NO	591-27-5	1298.61			60.7	NIST14
385	Sarcosylsarcosine, N-methoxycarbonyl-, heptyl ester	C15H28N2O5	1000314-83-0	1299.86			66.4	NIST14
386	Tridecane	C13H28	629-50-5	1300.00	1300.00	0.00	74.0	FingerprintingDB624
387	3,4-Dimethylphenol	C8H10O	95-65-8	1301.35	1302.07	0.71	60.3	FingerprintingDB624
388	4-Methyl-1,2-dihydronaphthalene	C11H12	4373-13-1	1309.40	1311.61	2.21	76.8	UCSD
389	Pentanedioic acid, 2-oxo-, dimethyl ester (CAS)	C7H10O5	13192-04-6	1310.40	1279.05	-31.34	49.4	UCSD
390	Nonanoic acid (CAS)	C9H18O2	112-05-0	1310.71	1323.04	12.33	54.4	UCSD
391	Cyclohexasiloxane, dodecamethyl-	C12H36O6Si6	540-97-6	1317.81			60.7	NIST14
392	Phenol,2,3,4-trimethyl	C9H12O	526-85-2	1320.41	1325.05	4.64	46.4	FingerprintingDB624
393	Succinic acid, 2,5-dimethylphenyl ethyl ester	C14H18O4	1000329-86-0	1324.09			56.4	NIST14
394	Naphthalene, 1,2-dihydro-3-methyl- (CAS)	C11H12	2717-44-4	1331.73	1322.44	-9.29	65.5	UCSD
395	1,4:3,6-Dianhydro-.alpha.-d-glucopyranose	C6H8O4	1000098-14-8	1332.95			77.8	NIST14
396	4-isopropylphenol	C9H12O	99-89-8	1334.55	1340.26	5.71	47.4	FingerprintingDB624
397	Divinylmethyl(chloromethyl)silane	C6H11ClSi	25202-02-2	1338.12			56.1	NIST14
398	1H-Pyrrole-2,5-dione, 3-ethyl-4-methyl- (CAS)	C7H9NO2	20189-42-8	1339.08	1364.74	25.66	53.3	UCSD

- Identified with PMI Accurate Mass Library (n=7 out of 21)
- Suspect compound library (n=6 out of 21)
- NIST proposal (n=8 out of 21)

Level of Confidence for Commercial Library proposals



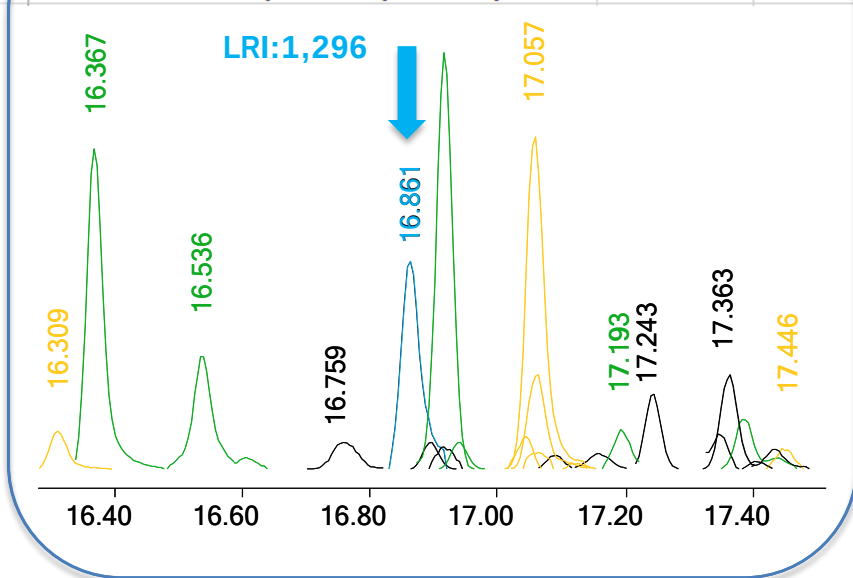
	A	B	C	D	E	F	G	H
1	Compound Name	Formula	CAS#	Component RI	Library RI	Δ RI	Match Factor	Library File
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385	Sarcosylsarcosine, N-methoxycarbonyl-, heptyl ester	C15H28N2O5	1000314-83-0	1299.86			66.4	NIST14
386								
387								
388								
389								
390								
391	Cyclohexasiloxane, dodecamethyl-	C12H36O6Si6	540-97-6	1317.81			60.7	NIST14
392								
393	Succinic acid, 2,5-dimethylphenyl ethyl ester	C14H18O4	1000329-86-0	1324.09			56.4	NIST14
394								
395	1,4:3,6-Dianhydro-.alpha.-d-glucopyranose	C6H8O4	1000098-14-8	1332.95			77.8	NIST14
396								
397	Divinylmethyl(chloromethyl)silane	C6H11ClSi	25202-02-2	1338.12			56.1	NIST14
398								

Pred LRI Rapid Miner	Pred LRI Chrom Genius	
1627.9	1776.4	Very different $\Delta = 412$
1164.0	1253.9	In agreement with exp $\Delta = 87$
1502.9	1477.3	Very different $\Delta = 191$
2304.4	2279.2	Very different $\Delta = 992$
4054.1	1635.9	Very different $\Delta = 1677$
1914.4	1934.4	Very different $\Delta = 600$
1354.0	1551.6	In agreement with exp $\Delta = 120$
990.8	943.7	Very different $\Delta = 371$

The identification confidence is low to medium with a commercial library search
 Filter out false positive results proposed by existing MS libraries

Case Study: Compound with a Poor Match Factor

	A	B	C	D	E	F	G	H
1	Compound Name	Formula	CAS#	Component RI	Library RI	Δ RI	Match Factor	Library File
383	Butanoic acid, 3-methyl-, 3-methyl-3-butenyl ester	C10H18O2	54410-94-5	1296.49			65.5	NIST14



Pred LRI
Rapid Miner 1164.0
Chrom Genius 1253.9
In agreement with exp $\Delta = 87$

LRI prediction

Data Acquisition
CI full scan MS
Targeted MS/MS

MetFrag
in silico
fragmentation software

Data Acquisition
EI full scan MS

CFM-ID
in silico
fragmentation software

Candidates

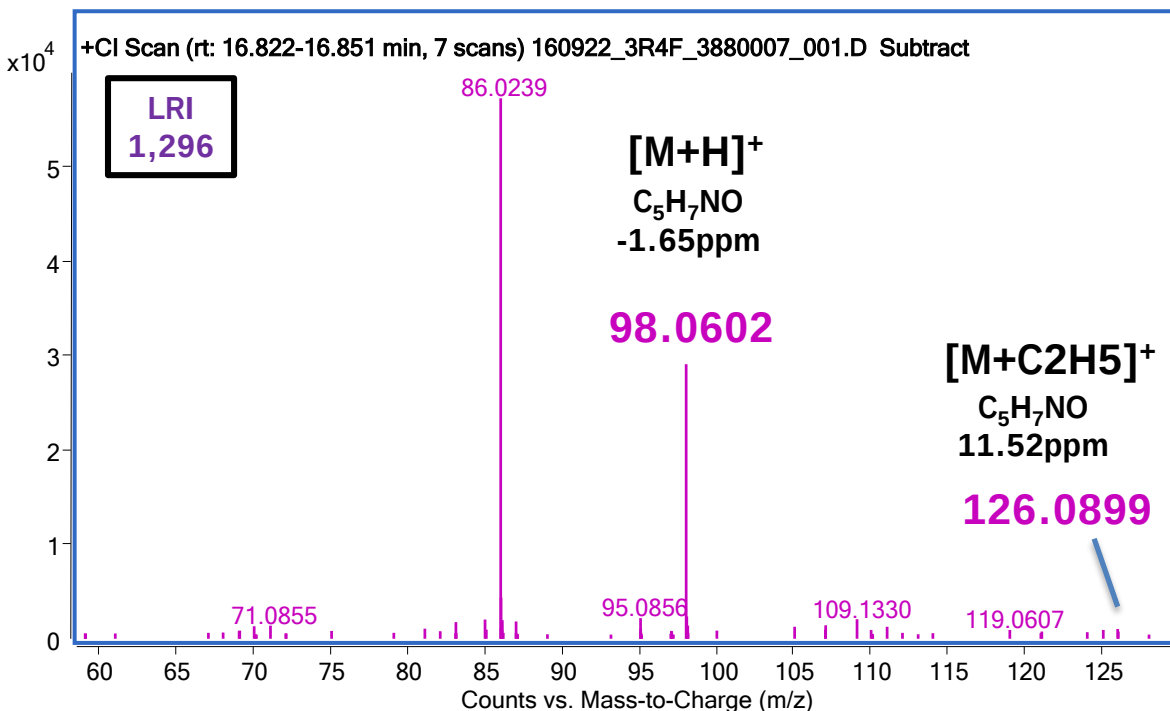
Final Score

Purchase of reference
standards if available

Additional MS experiment are needed
and alternative approaches need to
be implemented

GC-HR-MS in Chemical Ionization Mode & MS/MS

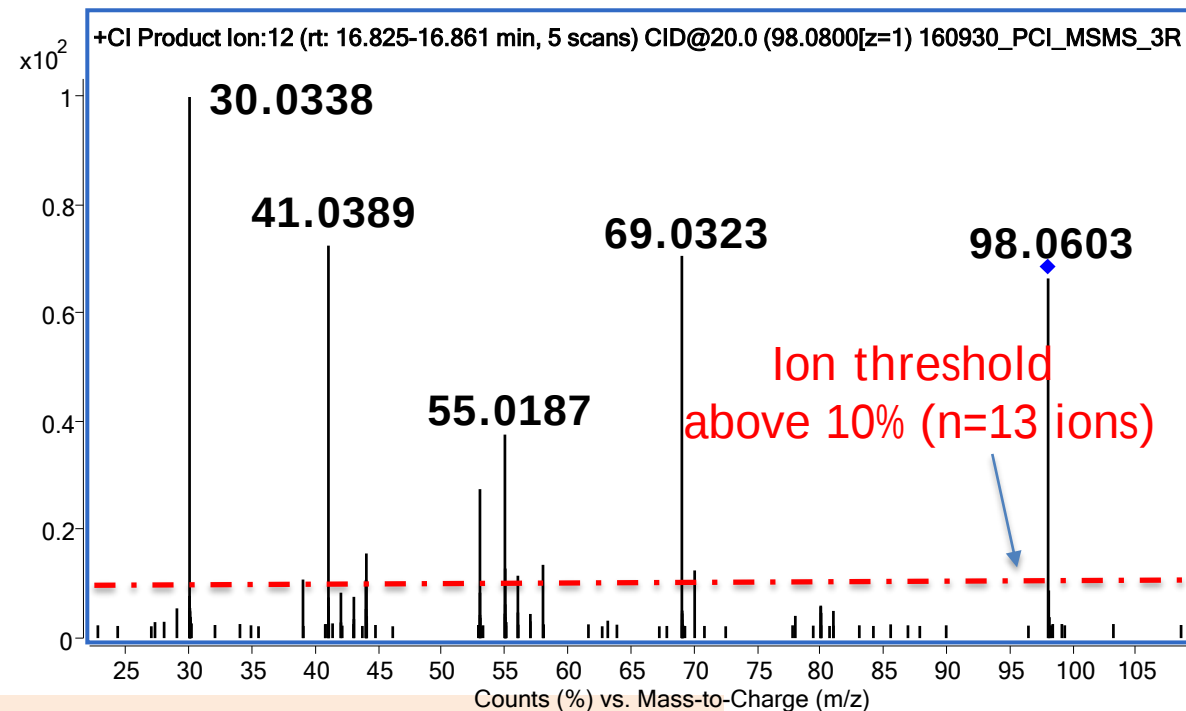
GC-HR-MS (Full Scan MS) Positive Chemical Ionization (PCI)



Determination of
elemental formula
(adduct ion species)

M: C₅H₇NO

GC-HR-MS (Full Scan MS/MS) PCI data acquisition CID of 98.08004



MS/MS data processed using
a larger chemical database
with *in silico* predicted
fragmentation software

Compound Identification with *In Silico* Fragmentation Software

MetFrag
In silico fragmentation for computer assisted identification of metabolite mass spectra

Database Settings

Database: Parent Ion:

Neutral Mass: Search ppm:

Formula:

Identifiers:

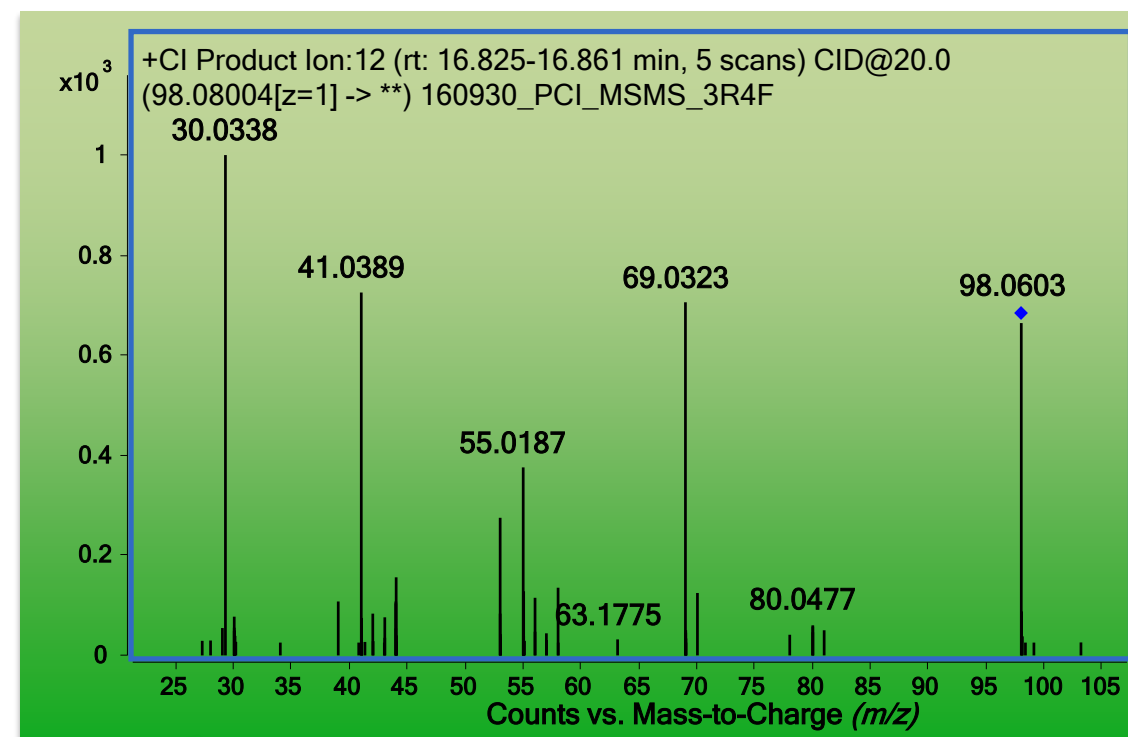
Fragmentation Settings & Processing

Mzppm:

Mzabs:

Mode:

MS/MS Peak list



C. Ruttkies, E.L. Schymanski, S. Wolf, J. Hollender, S. Neumann J.
Chemoinform. **2016**, 8, 3, 16-115.

Metfrag Results

Results

Download Results

Filter Candidates by explained MS/MS Peaks

MS/MS Peaks

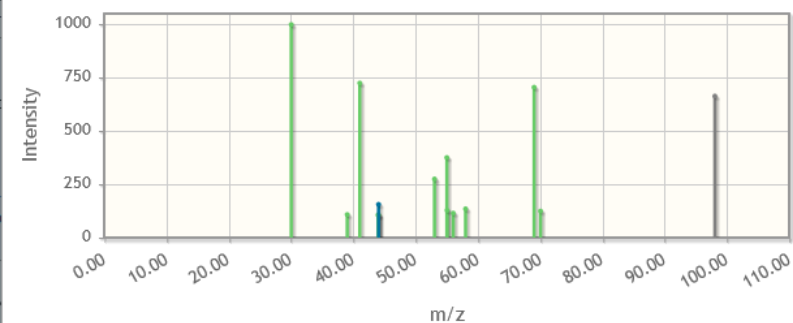
Filter Candidates

#	Molecule	Identifier	Mass	Formula	FinalScore	Details
1	 3-Pyrrolin-2-one, 1-methyl-	466735 InChIKeyBlock1 = VHGGRTWHRJRQKU	97.05277	C ₅ H ₇ NO	1.0	Peaks: 12 / 13 Fragments Scores Download
2	 1-Methyl-1,3-dihydro-2H-pyrrol-2-one	4483194 InChIKeyBlock1 = VIETUFSZPCIVQL	97.05277	C ₅ H ₇ NO	0.9913	Peaks: 12 / 13 Fragments Scores Download
3	 2-Methyl-2-azabicyclo[2.1.0]pentan-3-one	45531574 InChIKeyBlock1 = SYIPZAVNRAIEV	97.05277	C ₅ H ₇ NO	0.9758	Peaks: 12 / 13 Fragments Scores Download
4	 2-methyl-2-oxo-1,3-dihydro-2H-pyrrol-3-ylidene	32989542 InChIKeyBlock1 =	97.05277	C ₅ H ₇ NO	0.9589	Peaks: 10 / 13 Fragments Scores

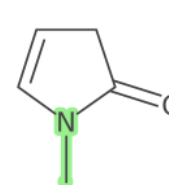
Fragments View

Select area to zoom in. Double click to return.
Click on apex of explained peak to select fragment.

matched
not matched
excluded

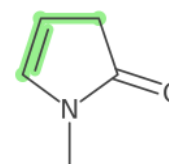


Fragments



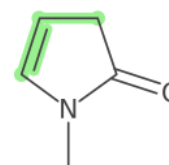
Fragment 1

Peak m/z: 30.0338
Fragment Mass: 30.03383 Da
Fragment Formula: [CH₃N]⁺H⁺



Fragment 2

Peak m/z: 39.0234
Fragment Mass: 39.02293 Da
Fragment Formula: [C₃H₄-H]⁺



Fragment 3

Peak m/z: 41.0389
Fragment Mass: 41.03858 Da
Fragment Formula: [C₃H₄]⁺H⁺

Compound Identification with Fragmentation Modeling Software

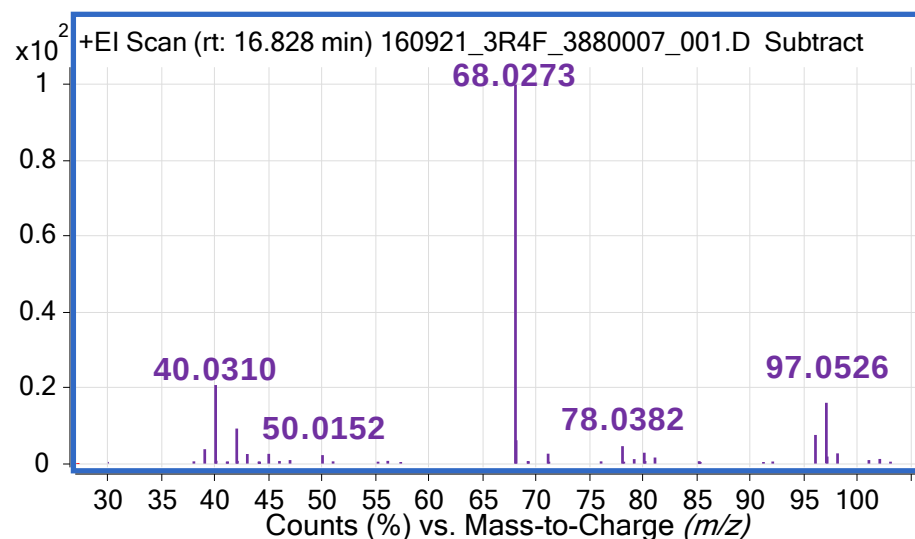
MetFrag
In silico fragmentation for computer assisted identification of metabolite mass spectra

Database Settings

Database: ChemSpider

4956336 InChI=1/C5H7NO/c7-5-3-1-2-4-6-5/h1,3H,2,4H2,(H,6,7)
10681140 InChI=1/C5H7NO/c7-5-1-3-6-4-2-5/h1,3,6H,2,4H2
21377662 InChI=1/C5H7NO/c6-4-1-2-5(7)3-4/h1-2,4H,3,6H2
.
14294945 InChI=1/C5H7NO/c7-5-4-1-3(4)2-6-5/h3-4H,1-2H2,(H,6,7)
13986409 InChI=1/C5H7NO/c6-5-3-1-2-4-7-5/h1-5H,6H2
21786394 InChI=1/C5H7NO/c1-3-4-5(7)6-2/h1H,4H2,2H3,(H,6,7)

178 Candidates Download Candidates



CFM-ID
Competitive Fragmentation Modeling for Metabolite Identification

Compound Identification

Determines the compounds that most closely match to a given spectra. The spectra for each candidate compound (in the provided list) are predicted using a pre-trained model and compared to the input spectra. The top candidates are ranked according to how closely they match and returned in a list.

Candidate Compounds OR

The candidates should be represented as a list of compounds in the format 'ID SMILES_or_InChI' on each line. The list can have a maximum of 100 compounds. Maximum individual compound size is 200 atoms. [Load an example.](#)

Input Text

Spectra Type: EI
Ion Mode: Positive
Spectra: OR

The spectra should be represented as a list of peaks with the format 'mass intensity' on each line, and can be entered directly into the corresponding energy level boxes below. Multiple energy levels are optional; only one is required.

Input Spectra Text
70eV Energy

Number of Results:
The number of results to return. Leave blank to return all results.

Mass Tolerance: Da
The mass tolerance to use when matching peaks within the spectrum comparison.

Scoring Function:

The EI spectra of each candidate are predicted using CFM-ID

CFM-ID Results

Results: [Download](#)

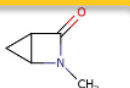
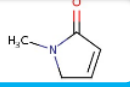
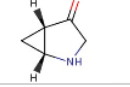
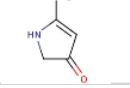
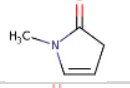
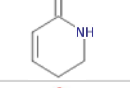
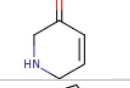
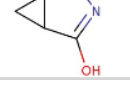
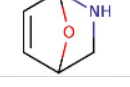
Input spectra are shown below in blue. If a database was queried, candidate spectra are overlayed on top for comparison. The top ranking candidate spectra is shown by default; to compare other database candidates use the "Compare" buttons on the list of ranked candidate compounds that follow the spectra.

Candidate Rankings

Rank	Score	Structure	ID	Chemical Formula	InChI/SMILES	Compare
1	0.27059567		45531574	C5H7NO	InChI=1/C5H7NO/c1-6-4-2-3(4)5(6)/h3-4H,2H2,1H3	Current
2	0.23287233		32990381	C5H7NO	InChI=1/C5H7NO/c7-5-2-6-4-1-3(4)5/h3-4,6H,1-2H2/t3-,4+/m0/s1	Compare
3	0.21118675		10340193	C5H7NO	InChI=1/C5H7NO/c1-4-2-5(7)3-6-4/h2,6H,3H2,1H3	Compare
4	0.19719996		27268587	C5H7NO	InChI=1/C5H7NO/c1-4-2-5(6)7-3-4/h2-3H,6H2,1H3	Compare
5	0.19212561		466735	C5H7NO	InChI=1/C5H7NO/c1-6-4-2-3-5(6)7/h2-3H,4H2,1H3	Compare
6	0.19070942		4956336	C5H7NO	InChI=1/C5H7NO/c7-5-3-1-2-4-6-5/h1,3H,2,4H2,(H,6,7)	Compare
7	0.1824996		14395305	C5H7NO	InChI=1/C5H7NO/c7-5-2-1-3-6-4-5/h1-2,6H,3-4H2	Compare



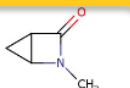
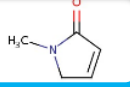
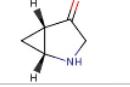
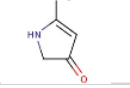
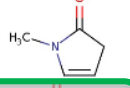
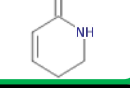
Final Score (MetFrag / CFM-ID)

InChI	CompoundName	Molecular Formula	Identifier	Structure	Rank MetFrag PCI	Score MetFrag PCI	Rank CFM-ID	Score CFM-ID
c1-6-4-2-3(4)5(6)7/h3-4H,2H2,1H3	2-Methyl-2-azabicyclo[2.1.0]pentan-3-one	C5H7NO	45531574		3	0.9758	1	0.271
c1-6-4-2-3-5(6)7/h2-3H,4H2,1H3	3-Pyrrolin-2-one, 1-methyl-	C5H7NO	466735		1	1.0	5	0.192
c7-5-2-6-4-1-3(4)5/h3-4,6H,1-2H2/t3-,4+/m0/s1	(1R,5S)-2-Azabicyclo[3.1.0]hexan-4-one	C5H7NO	32990381		16	0.9089	2	0.233
c1-4-2-5(7)3-6-4/h2,6H,3H2,1H3	5-Methyl-1,2-dihydro-3H-pyrrol-3-one	C5H7NO	10340193		15	0.915	3	0.211
c1-6-4-2-3-5(6)7/h2,4H,3H2,1H3	1-Methyl-1,3-dihydro-2H-pyrrol-2-one	C5H7NO	4483194		2	0.9913	11	0.167
c7-5-3-1-2-4-6-5/h1,3H,2,4H2,(H,6,7)	5,6-Dihydro-2(1H)-pyridinone	C5H7NO	4956336		5	0.9354	6	0.191
c7-5-2-1-3-6-4-5/h1-2,6H,3-4H2	1,6-Dihydro-3(2H)-pyridinone	C5H7NO	14395305		8	0.9294	7	0.182
c7-5-4-1-3(4)2-6-5/h3-4H,1-2H2,(H,6,7)	3-Azabicyclo[3.1.0]hexan-2-one	C5H7NO	14294945		12	0.9173	8	0.182
c1-2-5-6-3-4(1)7-5/h1-2,4-6H,3H2	7-Oxa-2-azabicyclo[2.2.1]hept-5-ene	C5H7NO	29542634		22	0.8925	9	0.181

$$\text{Combined Score} = (\text{MetFrag} \times 0.66 + \text{CFM-ID}_{\text{Norm}} \times 0.33)$$

Final Score (MetFrag / CFM-ID / LRI Predictions)

Unknown Peak
LRI_{exp}: 1,296

InChI	CompoundName	Molecular Formula	Identifier	Structure	Rank MetFrag PCI	Score MetFrag PCI	Rank CFM-ID	Score CFM-ID	Score MetFrag & CFM-ID	LRI RM	LRI CG	Final Score
c1-6-4-2-3(4)5(6)7/h3-4H,2H2,1H3	2-Methyl-2-azabicyclo[2.1.0]pentan-3-one	C5H7NO	45531574		3	0.9758	1	0.271	0.974	962.51	1036.66	0.65
c1-6-4-2-3-5(6)7/h2-3H,4H2,1H3	3-Pyrrolin-2-one, 1-methyl-	C5H7NO	466735		1	1.0	5	0.192	0.894	995.41	1168.08	0.66
c7-5-2-6-4-1-3(4)5/h3-4,6H,1-2H2/t3-,4+/m0/s1	(1R,5S)-2-Azabicyclo[3.1.0]hexan-4-one	C5H7NO	32990381		16	0.9089	2	0.233	0.884	1126.04	1015.88	0.64
c1-4-2-5(7)3-6-4/h2,6H,3H2,1H3	5-Methyl-1,2-dihydro-3H-pyrrol-3-one	C5H7NO	10340193		15	0.915	3	0.211	0.861	1057.08	918.43	0.56
c1-6-4-2-3-5(6)7/h2,4H,3H2,1H3	1-Methyl-1,3-dihydro-2H-pyrrol-2-one	C5H7NO	4483194		2	0.9913	11	0.167	0.858	994.94	1083.00	0.60
c7-5-3-1-2-4-6-5/h1,3H,2,4H2,(H,6,7)	5,6-Dihydro-2(1H)-pyridinone	C5H7NO	4956336		5	0.9354	6	0.191	0.850	1110.21	1304.80	0.74
c7-5-2-1-3-6-4-5/h1-2,6H,3-4H2									0.836	1170.32	1049.83	0.64
c7-5-4-1-3(4)2-6-5/h3-4H,1-2H2,(H,6,7)	3-								0.827	1041.30	1303.53	0.69
c1-2-5-6-3-4(1)7-5/h1-2,4-6H,3H2	7-Ox								0.810	1098.64	893.42	0.54

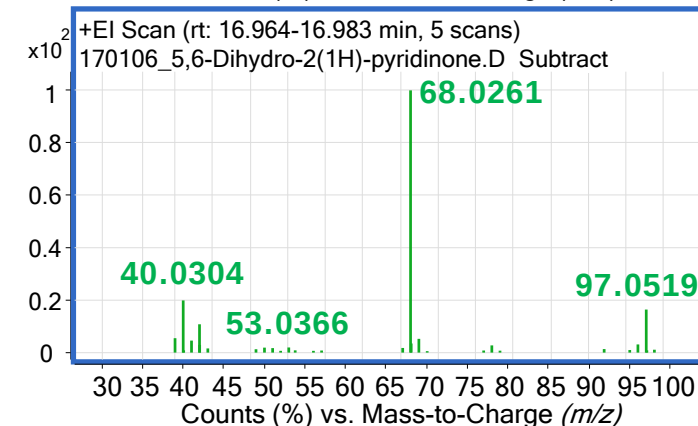
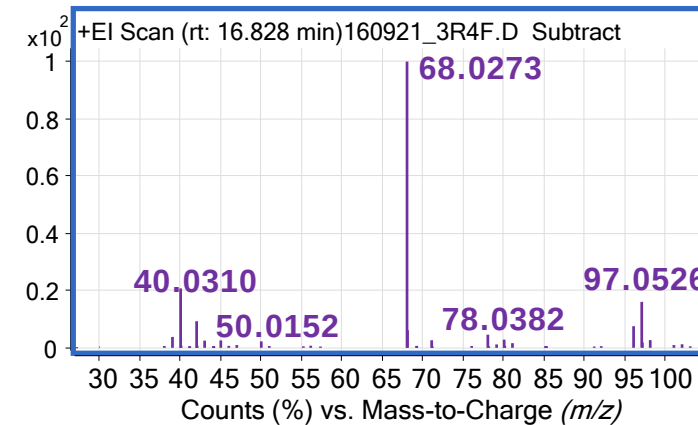
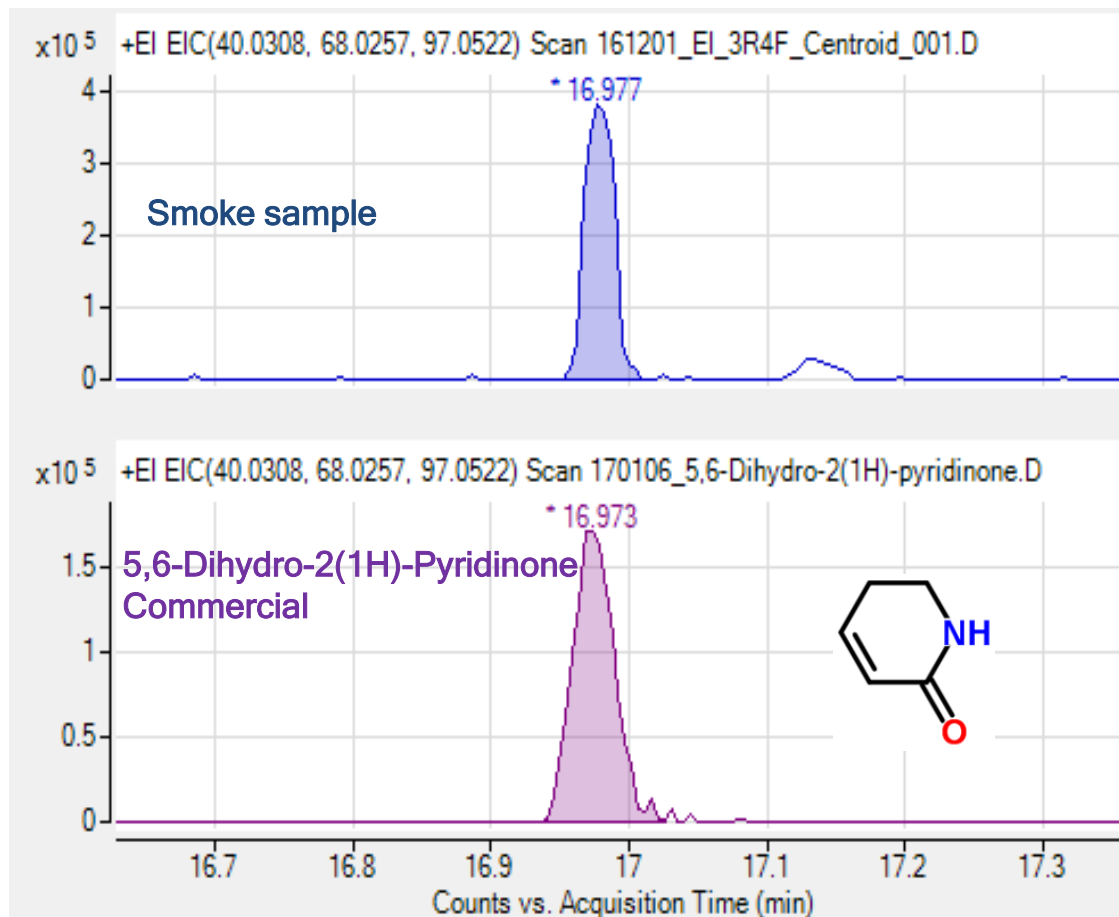
1) We use Inchi information to draw the compound and LRI values were predicted for all 178 proposals

2) Final SCORE was calculated using:

- ✓ MetFrag Score
- ✓ CFM-ID Score
- ✓ LRI_{exp.} against LRI_{RM}
- ✓ LRI_{exp.} against LRI_{CG} ...

$$\text{Final}_{\text{Score}} = \frac{(\text{MetFrag} \times 0.66 + \text{CFM-ID}_{\text{Norm}} \times 0.33) \times \text{LRI}_{\text{exp}} \times \text{LRI}_{\text{exp}}}{\left[\text{LRI}_{\text{exp}} + \sqrt{(\text{LRI}_{\text{CG}} - \text{LRI}_{\text{exp}})^2} \right] \times \left[\text{LRI}_{\text{exp}} + \sqrt{(\text{LRI}_{\text{RM}} - \text{LRI}_{\text{exp}})^2} \right]}$$

Final Confirmation



Similar EI Mass Spectra

Same RT

5,6-Dihydro-2(1H)-Pyridinone ✓

Bibliography

NIH > NLM > National Center for Biotechnology Information

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Search Compounds

Compound Summary for CID 6453994

Download Share Help

5,6-Dihydropyridin-2(1H)-one

Cite this Record



PubChem CID:	6453994
Chemical Names:	5,6-Dihydropyridin-2(1H)-one; 6052-73-9; 5,6-Dihydro-2(1H)-pyridone; 5,6-Dihydro-2(1H)-pyridinone; 5,6-DIHYDRO-1H-PYRIDIN-2-ONE; 2(1H)-Pyridone, 5,6-dihydro- More...
Molecular Formula:	C ₅ H ₇ NO
Molecular Weight:	97.117 g/mol
InChI Key:	OXRHRHYRQWIIHV-UHFFFAOYSA-N
Substance Registry:	FDA UNII

PUBCHEM > COMPOUND > 5,6-DIHYDROPYRIDIN-2(1H)-ONE

Modify Date: 2017-01-07; Create Date: 2006-04-29

Contents

7.1 Depositor Provided PubMed Citations

Download

1 to 3 of 3			
PMID	Date	Title	Journal
23434420	2013-04-01	Anti-HBV active constituents from Piper longum.	Bioorganic & medicinal chemistry letters
20481450	2010-06-18	Synthesis of densely substituted trans-configured 4-acylated piperidine-2,4-diones as 3:1 adducts of imines and ketenes.	The Journal of organic chemistry
17869121	2007-12-01	Synthesis and biological evaluation of non-peptide alpha(v)beta(3)/alpha(5)beta(1) integrin dual antagonists containing 5, 6-dihydropyridin-2-one scaffolds.	Bioorganic & medicinal chemistry

Smoke Composition. An Extensive Investigation of the Water-Soluble Portion of Cigarette Smoke

Joseph N. Schumacher,* Charles R. Green, Freddie W. Best, and Marjorie P. Newell

Research Department, R. J. Reynolds Tobacco Company, Winston-Salem, North Carolina 27102.

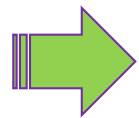
310 J. Agric. Food Chem., Vol. 25, No. 2, 1977

- This compound was reported first from cigarette smoke by Schumacher et *al.* in 1977
- NOT registered in NIST 14 NOR in Wiley 11
- Strengthen the need to implement *in silico* predictions when dealing with untargeted analysis.

Conclusion



Existing MS libraries are not exhaustive; the implementation of chemoinformatic tools is needed to postulate compounds that are not registered in any library



The combined use of MetFrag and CFM-ID software has been demonstrated to be a good complementary tool to propose reliable compound hits



Addition of LRI predictions demonstrated the ability to correctly rank putative hits proposed by *in silico* fragmentation software



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Thank You

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Philip Morris International R&D, Philip Morris Products S.A.,
(part of Philip Morris International group of companies)