



PMI RESEARCH & DEVELOPMENT

Computational Platform for Compound Identification

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Philip Morris International, R&D

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ACD/Labs Symposium on Laboratory Intelligence (EUM), Neuchatel Switzerland

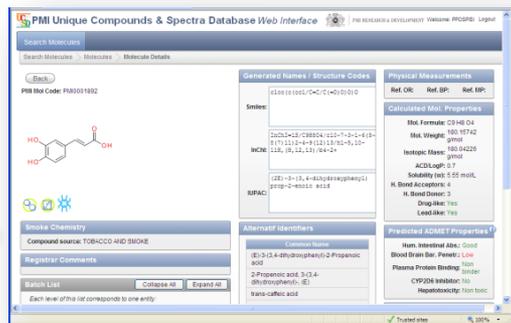
Introduction

Chemoinformatics challenges at PMI:

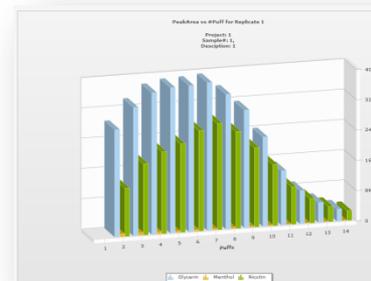
- Compound identification from complex matrices
- Efficient and/or automatized compound registration (including registration of mixtures and stereochemical isomers)
- Managing spectral data
- Associating toxicity data to compounds and mixtures
- Inserting chemical data in corporate Scientific Data Warehouse
- Reporting chemical data in a relevant way
- Building R&D chemoinformatics platform using Rapid Development Tools



PMI Chemoinformatics platform



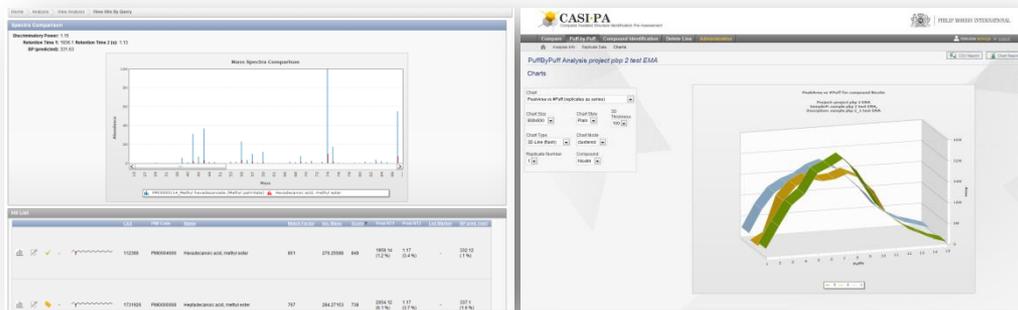
1. Unique Compounds & Spectra Database



3. Chemoinformatics Knowledge Base



Scientific Data Warehouse



2. Computer Assisted Structure Identification



1. Unique Compounds & Spectra Database



1. Unique Compounds & Spectra Database



3. Chemoinformatics Knowledge Base

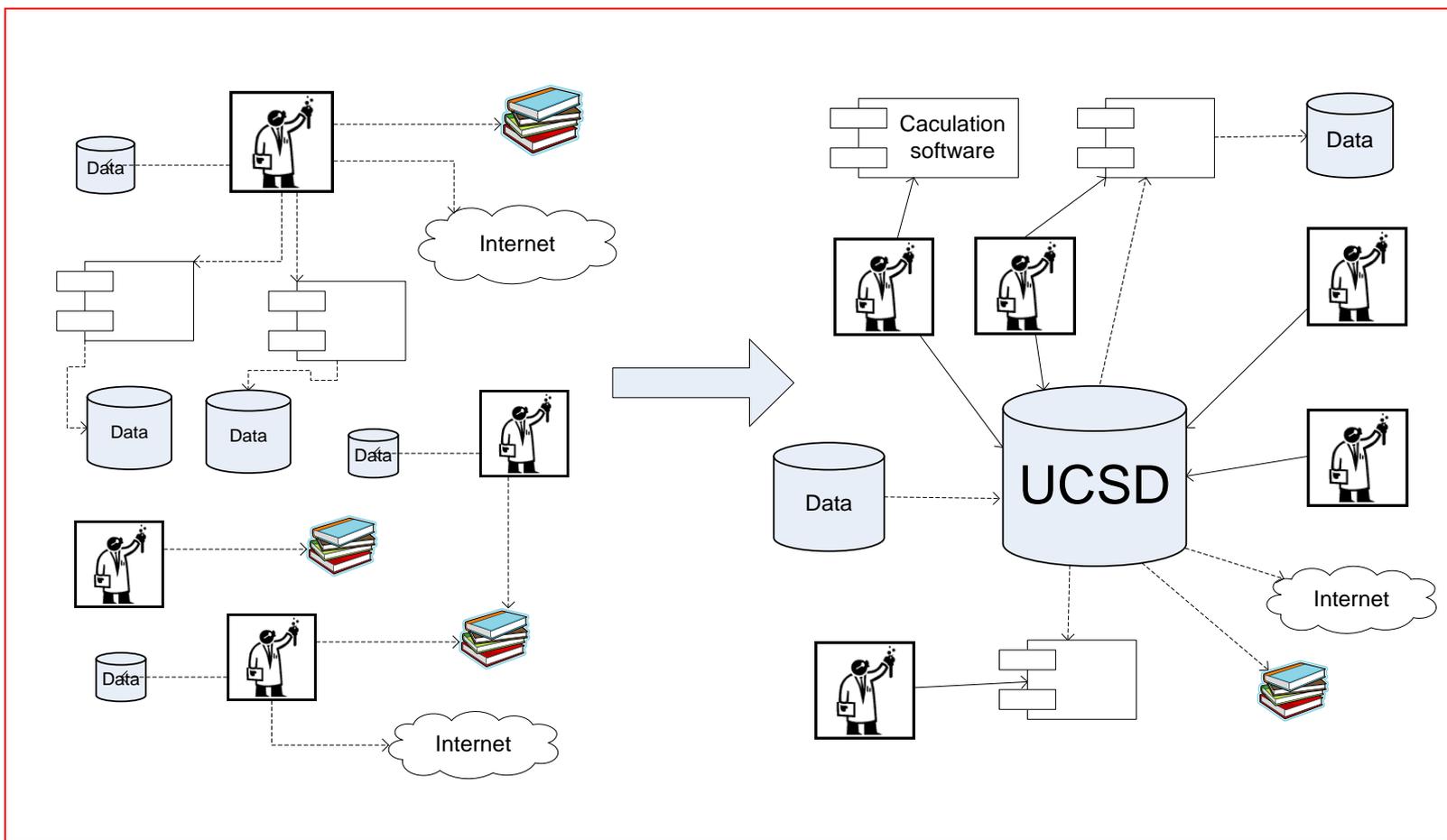


Scientific Data Warehouse

2. Computer Assisted Structure Identification



Concept of UCSD

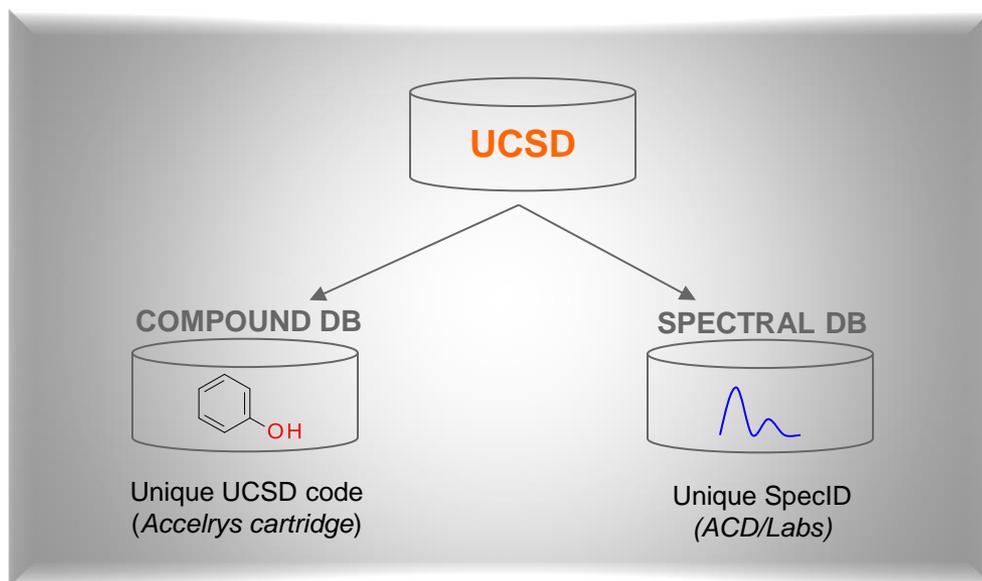


Unique Compound and Spectra Database: To assemble R&D chemical information into a central repository of chemical substances and analytical spectra.



Concept of UCSD

- UCSD is an internal database with no external access
- Accelrys chemical cartridge with enhanced stereochemistry
- ACD/Labs for the analytical spectra part
- Web interface developed in Oracle Application Express and automatic processes in Accelrys Pipeline Pilot



UCSD Roles



1. Submitter

A chemist who can **search**, **view** data and also **submit** molecules and spectra.



2. Registrar

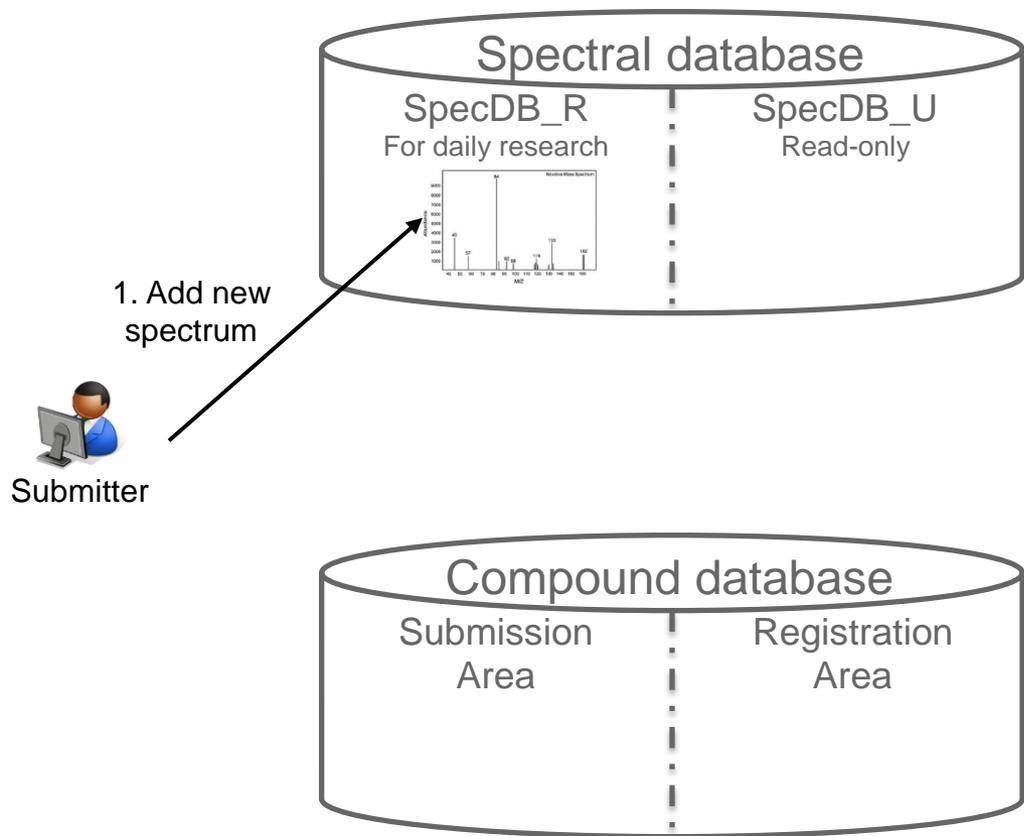
A chemoinformatician that can **search**, **view**, **submit** and also check and **register** data.



Viewer

User can search and **view** registered data (print, export, list creation)





ACD/ChemFolder Enterprise module

ACD/ChemFolder Enterprise: Database Window - [emartin is now connected to CHEMBOOK_S... on rd-acd-chemfolder.app.pmi <Default>]

Database View Record Search Lists Plates Options ACD/Labs Help

LOCAL REMOTE

Tile Table Default with User Data CFE_main Update current record Add spectrum to record ImportCsv ExportToIDX ImportMZMine

Short File Name: _1_MASS SPECTRUM
Count: 31

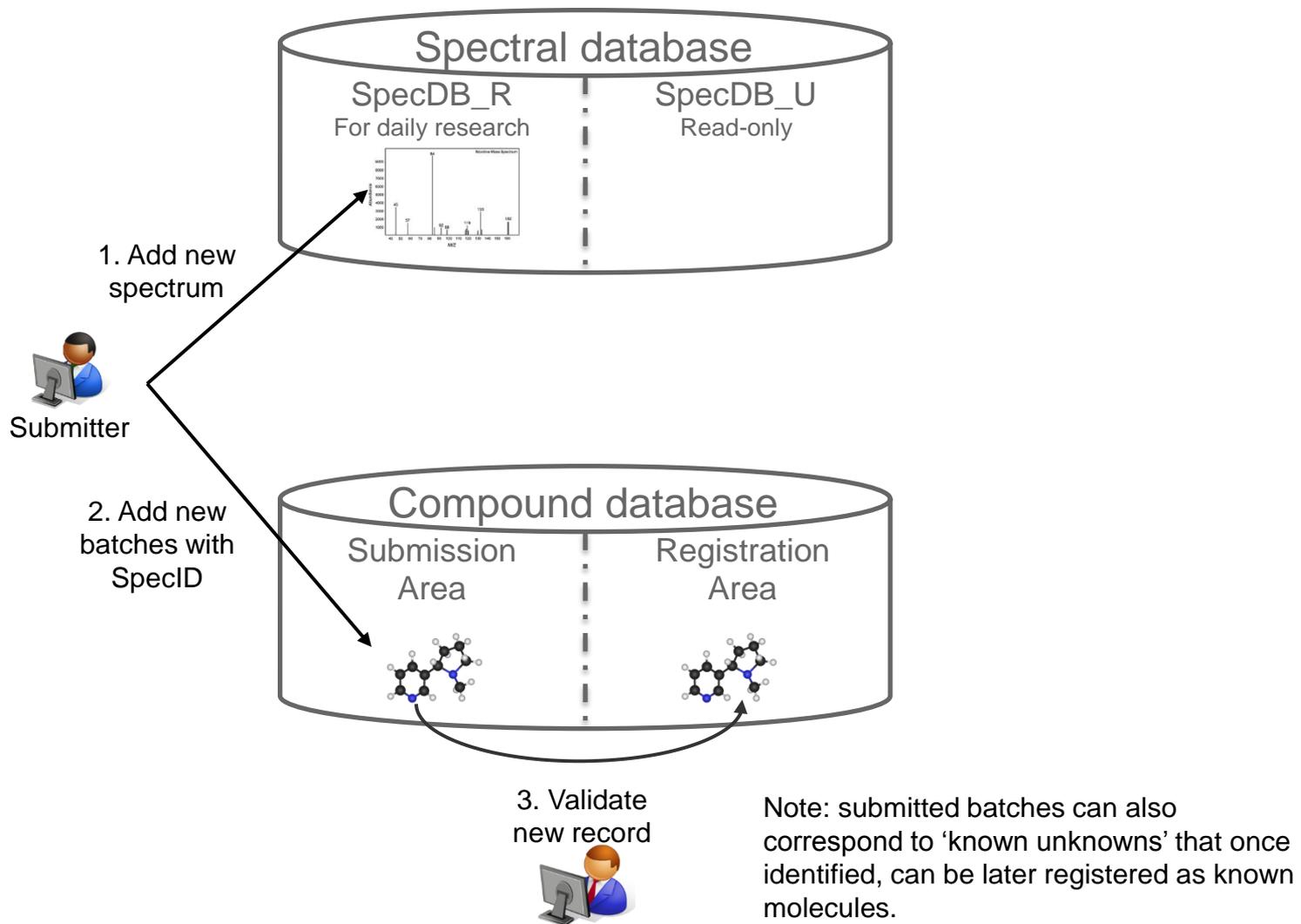
Name	Value
CAS Name	ISOPRENE_2-METHYL-1,3-BUTADIENE
CAS Registry Num	78795
Count	31
Data Type	MASS SPECTRUM
Estimated Molecul	68.000
Molecular Formula	CSH8
Molecular Weight	68
Spectrum Title	PYRO_PASS entry 126
TIC	433.71
Total Signal	433.67

Submitter	emartin	SpecID	SPR000000391
Scientist	Polier-Calame Josée	Project	Product Assessment
SDMS_ID		Study number	RLS-2012-VALID-108
Ionization type	EI	Analyzer type	single-quad
Adduct		Experimental mass	
Proposed element composition		Machine type	Pyro_PyR-GC-MS_8288
Theoretical monoisotopic mass		Product/Sample	
Theoretical monoisotopic mass of neutral molecule		Compound class	
Column name	Varian CP 5845 CP-SIL 8_Lenght 50m_ID 0.25mm_Film 0.4micron	Method name	PMI_RD_WK1_000387_Direct and indirect materials analysis by Pyrolysis-GC-MS
Retention time 1D	6.28	Retention time 2D	
Kovats index		Relative retention time	
Confidence	Reference	Copied in SpecDB_U	true

ID: 387 A: 254/272 B: 272 Single DB Owners: EMARTIN

1-ChemSketch 2-Database





Concept of Molecule, Substance and Batch Assignment

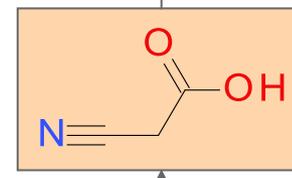
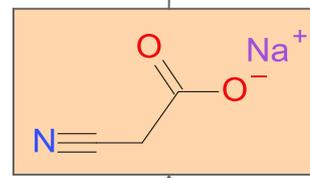
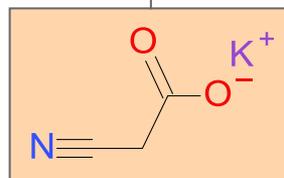
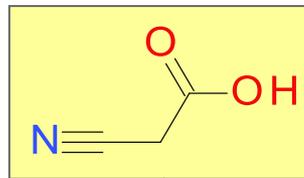
Unique means there is no redundancy, ensured by uniqueness of chemical structure and systematic and accurate registration process.

Molecule =
Neutral, unique
chemical entity

Substance =
Molecule + Salt

Batch =
Substance physically
present or project-
relevant as cited in
literature

Every molecule is
assigned a unique
company code:
e.g., **PMI01234567**

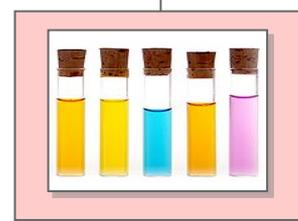
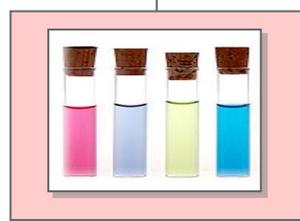


Substance ID = e.g.,

PMI01234567-A

PMI01234567-B

PMI01234567-C



Batch ID = e.g.,

BC000002152
BC000008641
BC000000560
BC000000320

BC000000122
BC000000154
BC000000176
BC000002504
BC000000894

BC000000121
BC000000158
BC000003174
BC000002598



UCSD Compound Part

The screenshot displays the web interface for the PMI Unique Compounds & Spectra Database. The main content area is divided into several panels:

- Registrars Comments:** A section for user comments.
- Batch List:** A tree view showing the hierarchy of batches. The selected batch is PM10000048, which contains sub-batches: PM10000048-A, BC000000055, BC000000056, BC000002843, BC000010355, and *BC000010489.
- List:** A table with columns for Name, FDA list, and Health Canada. It shows 1 - 2 entries.
- Generated Names / Structure Codes:** A table with columns for Smiles, InChI, and IUPAC. The values are:
 - Smiles: CC(=C)C=C
 - InChI: InChI=1S/CSH8/c1-4-5(2)/3/H4H,1-2H2,3H3
 - IUPAC: 2-methylbuta-1,3-diene
- Physical Measurements:** A table with columns for Ref. OR, Ref. BP, and Ref. MP.
- Calculated Mol. Properties:** A table with columns for Mol. Formula, Mol. Weight, Isotopic Mass, ACD/LogP, Solubility (w), H. Bond Acceptors, H. Bond Donor, Drug-like, and Lead-like. The values are:
 - Mol. Formula: C5H8
 - Mol. Weight: 68.11702 g/mol
 - Isotopic Mass: 68.0626 g/mol
 - ACD/LogP: 2.3
 - Solubility (w): 0.01 mol/L
 - H. Bond Acceptors: 0
 - H. Bond Donor: 0
 - Drug-like: Yes
 - Lead-like: Yes
- Predicted ADMET Properties:** A table with columns for Hum. Intestinal Abs., Blood Brain Bar. Penetr., Plasma Protein Binding, CYP2D6 inhibitor, and Hepatotoxicity. The values are:
 - Hum. Intestinal Abs.: Moderate
 - Blood Brain Bar. Penetr.: High
 - Plasma Protein Binding: Non binder
 - CYP2D6 inhibitor: No
 - Hepatotoxicity: Toxic
- Literature Data:** A section for literature references.
- Alternatif Identifiers:** A table with columns for Common Name, CAS, and IUPAC. The values are:
 - Common Name: 1,3-Butadiene, 2-methyl; 1,3-Butadiene, 2-methyl(isoprene); 2-METHYL-1,3-BUTADIENE; ISOPRENE; isoprene
 - CAS: 78-79-5
 - IUPAC: 1 - 1
- Chemical Classification:** A section for chemical classification.

• To each newly registered compound we developed automatic processes that:

- attach SMILES strings, InChI codes and IUPAC names

- calculate structure-derived physico-chemical and ADMET properties

⇒ Naming and calculation of compounds **after** registration (uniqueness check) reduces further ambiguity

See more in Martin et al., *Journal of Cheminformatics* 2012, 4:11



UCSD Compound Part

UCSD PMI Unique Compounds & Spectra Database Web Interface

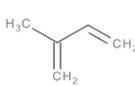
PMI RESEARCH & DEVELOPMENT Welcome: EMARTIN Logout

Search Molecules Submit New Molecule Administration

Search Molecules > Molecules > Molecule Details

Back

PMI Mol Code: PMI0000048



Generated Names / Structure Codes

Smiles: CC(=C)C=C

InChI: InChI=1S/CSH8/c1-4-5(2)/3/4H,1-2H2,3H3

IUPAC: 2-methylbuta-1,3-diene

Physical Measurements

Ref. OR: Ref. BP: Ref. MP:

Calculated Mol. Properties

Mol. Formula: C5H8
Mol. Weight: 68.11702 g/mol
Isotopic Mass: 68.0626 g/mol
ACD/LogP: 2.3
Solubility (w): 0.01 mol/L
H. Bond Acceptors: 0
H. Bond Donor: 0
Drug-like: Yes
Lead-like: Yes

Predicted ADMET Properties

Registrar Comments

Batch List

Each level of this list corresponds to one entity:
- molecule (neutral compound, e.g. PMI0000001)
- substance (molecule + counterion, hydrate e.g. PMI0000001-A)
- batches (physical or literature substances, e.g. BC0000014).

Click on a batch code to see its details.
Batches matched by the search query are marked with a *.

- PMI0000048
 - PMI0000048-A
 - BC000000055
 - BC000000056
 - BC000002843
 - BC000010355
 - * BC000010489

List

Name
FDA list
Health Canada

1 - 2

UCSD PMI Unique Compounds & Spectra Database Web Interface

PMI RESEARCH & DEVELOPMENT Welcome: EMARTIN Logout

Search Molecules Submit New Molecule Administration

Search Molecules > Molecules > Molecule Details > Substance & Batch

Back Substance

Details

Substance Code: PMI0000048-A

Salt	Coefficient
No salt	1

1 - 1

In UCSD a substance is equivalent to a molecule (which is always neutral in UCSD) with the additional information of the salts associated to the molecule.

Physical Measurements

Opt. rot.: Boil. point: Melt. point:
SpecID: SPR000000391

Calculated Properties

Mol. Weight: 68.117 g/mol Iso. Mass: 68.0626 g/mol
Mol. Formula: C5H8

Information (CAS and literature)

Batch

Archive Batch Edit Batch Structure

Details

Batch Code: BC000010489

Vial Barcode:

In UCSD each physical sample of a substance is registered as a separate batch.

Source Data

Project:

Scientist: Polier-Calame Josée Labbook ID:

Prod. date: Batch source: COMMERCIAL

Submitter: Bulk import Smoke:

Occurrence

AEROSOL

1 - 1

Comments

Stereo Comment:

Registrar comment:

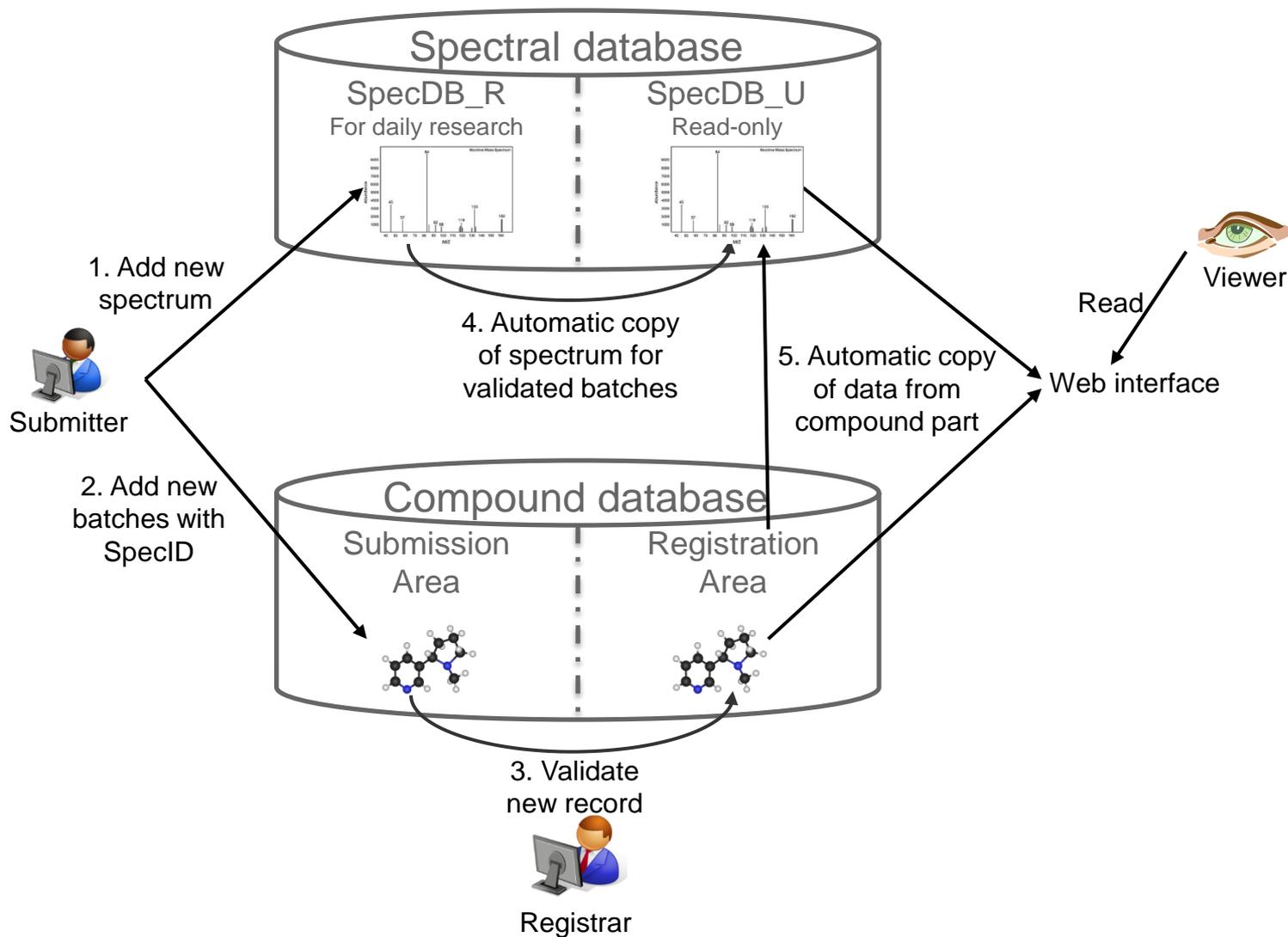
General Comments

PYRO_PASS entry 126

1 - 1

See more in Martin et al., *Journal of Cheminformatics* 2012, 4:11





ACD/ChemFolder Enterprise: Database Window - [emartin is now connected to CHEMBOOK_S... on rd-acd-chemfolder.app.pmi <Default>]

Database View Record Search Lists Plates Options ACD/Labs Help

LOCAL REMOTE [Icons]

File Table Default with User Data CFE_main2 Update current record Add spectrum to record ExportToJDX ExportToNIST Archive

Short File Name: _1_MASS SPECTRUM
Count: 31

C=CC=C

Submitter
emartin

Scientist
Polier-Calame Josée

Project

SDMS_ID

Study number
RLS-2012-VALID-108

Product/Sample

SpecID
SPR00000391

batchcode
BC000010489

Compound class

Machine type
Pyro_PYR-GC-MS_8288

Proposed element composition

Analyzer type
single-quad

Experimental mass

Theoretical monoisotopic mass of neutral molecule

Method name
PMI_RD_WK1_000387_Direct and indirect materials analysis by Pyrolysis-GC-MS

Retention Time 2D

Relative retention time

Name	Value
CAS Name	ISOPRENE_2-METHYL-1,3-BUTADIENE
CAS Registry Num	78795
Count	31
Data Type	MASS SPECTRUM
Estimated Molecul	68.000
Molecular Formula	CSH8
Molecular Weight	68
Spectrum Title	PYRO_PASS entry 126
TIC	433.71
Total Signal	433.67

ID: 253 A: 253/271 B: 271

Single DB

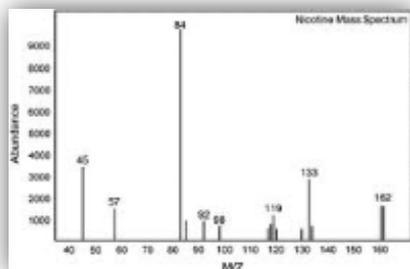
Owners: CHEMFOLDER_SCRIPT

1-ChemSketch 2-Database



In waiting for Spectrus

Search by spectra similarity is not possible in ChemFolder module → implementation of a script that copies each entry in Spectrum Database module



**ChemFolder
Enterprise**

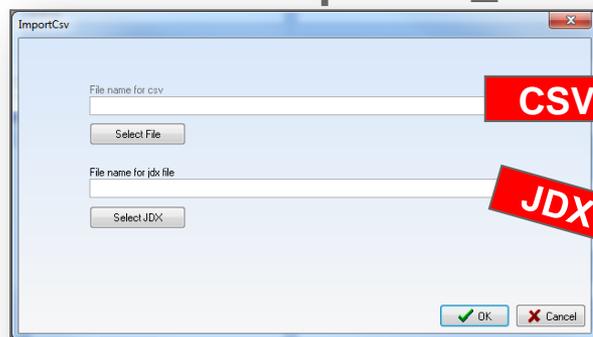


**Spectrum
Database**



Process for bulk import

INPUT for SpecDB_R



CSV file

JDX file

A	B	C	D	E	F	G	H
Title	Submitter	Scientist	Project	Confidence	Ionization type	Analyzer type	Pro
2-Nonanone-1,1,1,3,3-d5	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C9D	
Nitrobenzene-d5	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C6D	
O-cresol-d8	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C7D	

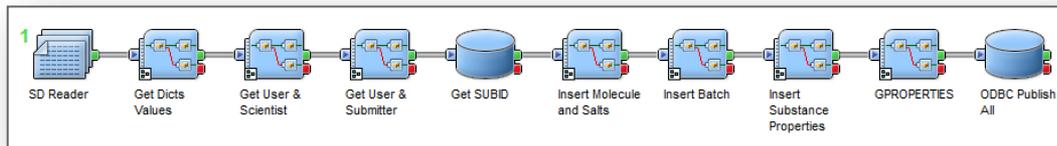
```
##TITLE= 2-Nonanone-1,1,1,3,3-d5
##DATA TYPE= MASS SPECTRUM
##MW= 147.1671489294
##MOLFORM=C9D5H13O
##CAS REGISTRY NO=1398065-76-3
##CAS NAME=2-Nonanone-1,1,1,3,3-d5
##NAMES=PMI0008955
##NPOINTS= 87
##XYDATA= (XY..XY)
30.0629, 0.996579903650516 31.0520, 1.03540117877503
##END=
##TITLE= Nitrobenzene-d5
##DATA TYPE= MASS SPECTRUM
##MW= 128.0634121394
##MOLFORM=C6D5NO2
##CAS REGISTRY NO=4165-60-0
```

OUTPUT

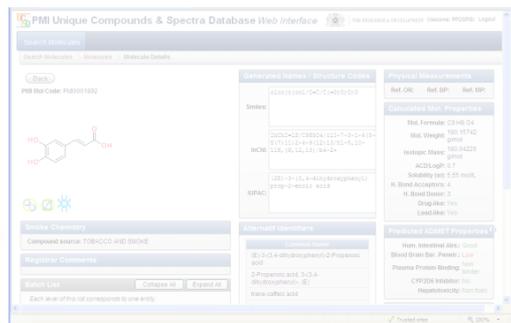
A	B	C	D	E	F	G	H	
Title	SpecID	Submitter	Scientist	Project	Confidence	Ionization type	Analyzer type	Pro
2-Nonanone-1,1,1,3,3-d5	SPR000000411	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C9D	
Nitrobenzene-d5	SPR000000412	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C6D	
O-cresol-d8	SPR000000413	emartin	Dossin Eric	Chemistry to Exposure Reference	EI	QTOF	C7D	



Merge and import in the compound part via Pipeline Pilot protocol



2. Computer Assisted Structure Identification



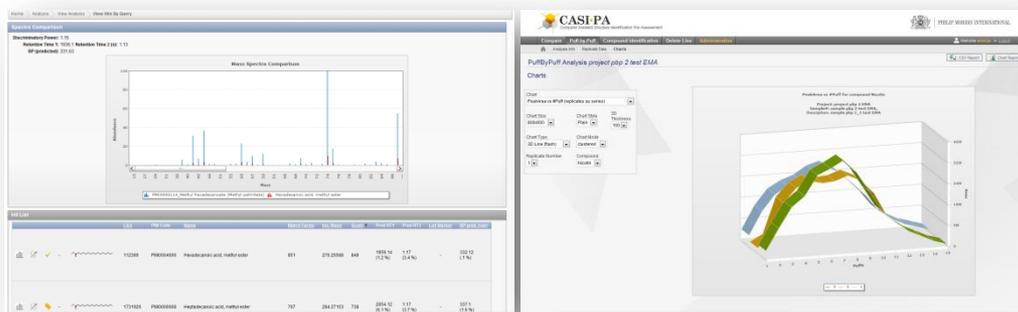
1. Unique Compounds & Spectra Database



3. Chemoinformatics Knowledge Base



Scientific Data Warehouse

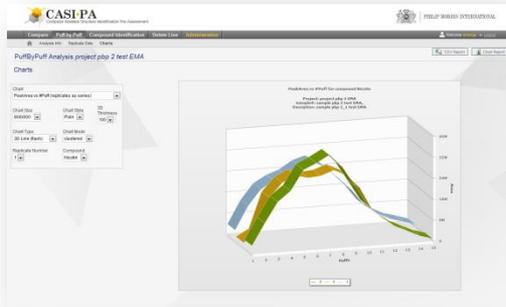


2. Computer Assisted Structure Identification



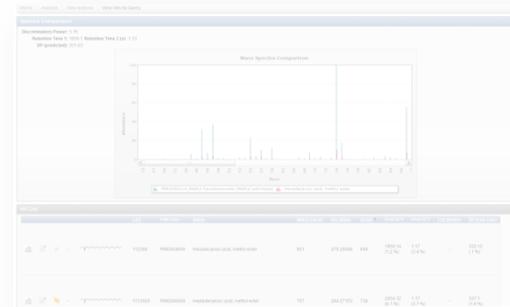
Computer Assisted Structure Identification

Our in-house developed Computer Assisted Structure Identification platform is constituted of two software:



Software 1 “CASI-PA”: assists the processing of MS data for given analyses:

- Compound Identification
- Comparison of 2 samples
- Puff-by-Puff

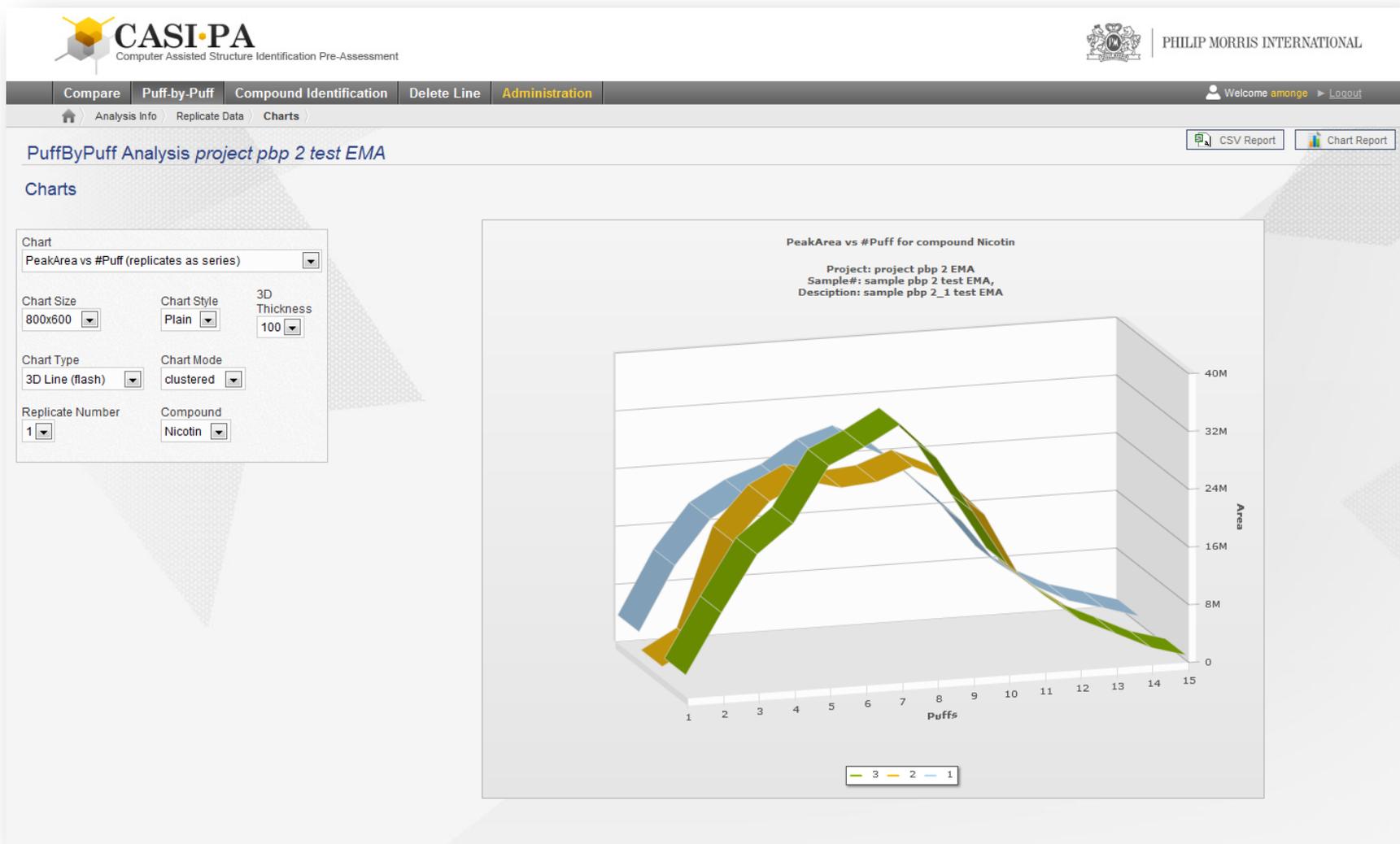


Automated non-target processing of MS data.



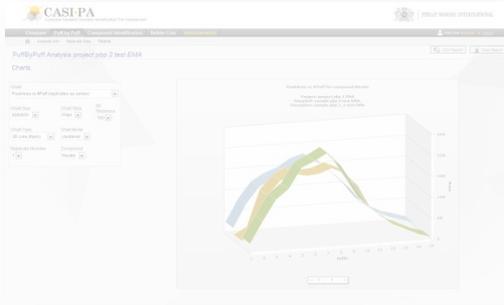
Puff-by-Puff Analysis of Cigarette Aerosols

Analysis of the concentration of given chemical compounds per puff.



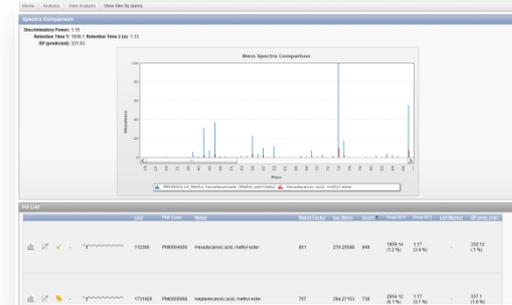
Computer Assisted Structure Identification

Our Computer Assisted Structure Identification platform is constituted of two software:



Help for the processing of MS data for given analyses:

- Compound Identification
- Puff-by-Puff
- Comparison of two products



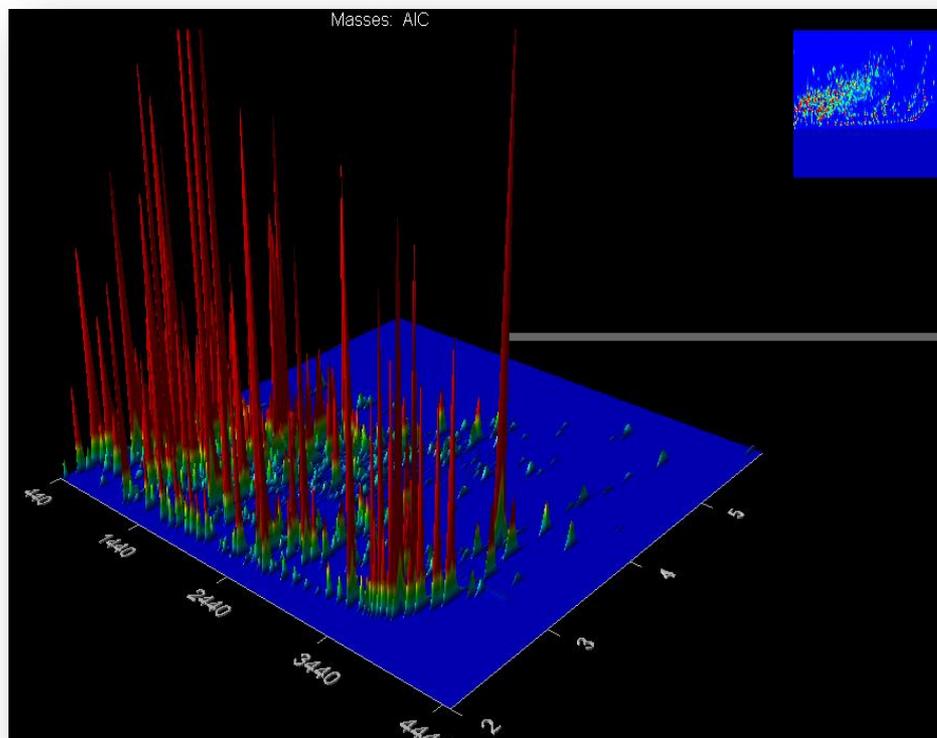
Software 2: Automated non-target processing of MS data.



Computer Assisted Structure Identification

Automated platform to accelerate and standardize identification of chemical structures of aerosol constituents with high confidence power.

Smoke of a conventional cigarette, measured by GCxGC-EI-TOF-MS



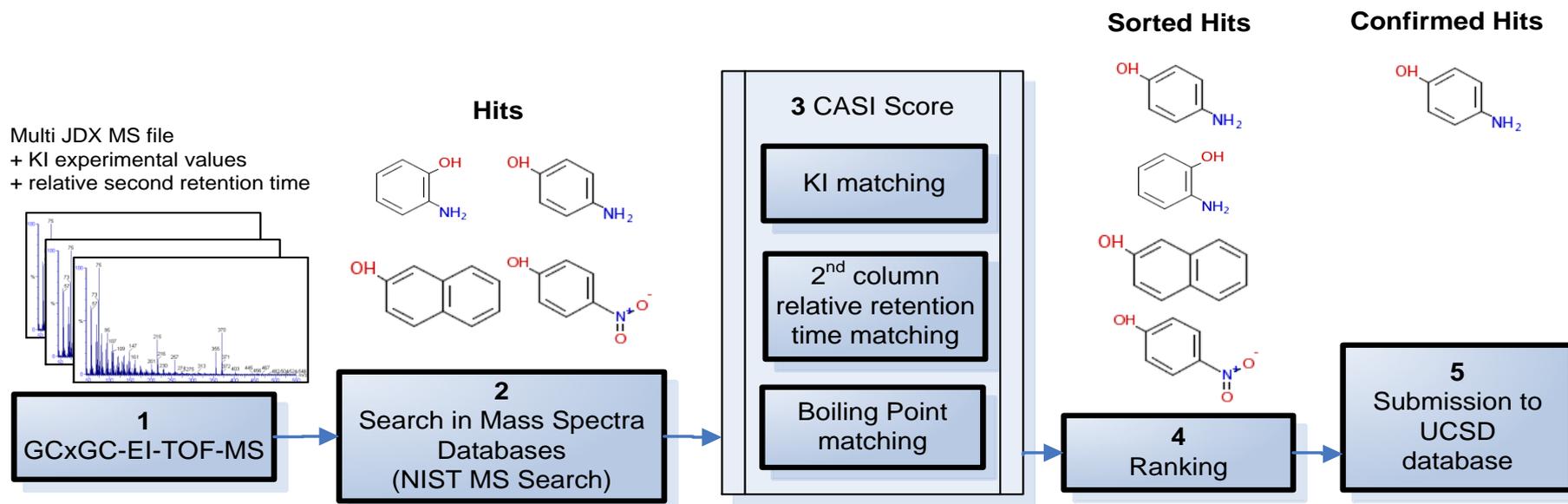
Compound?

A. Knorr et al., Anal. Chem. 2013, submitted for publication



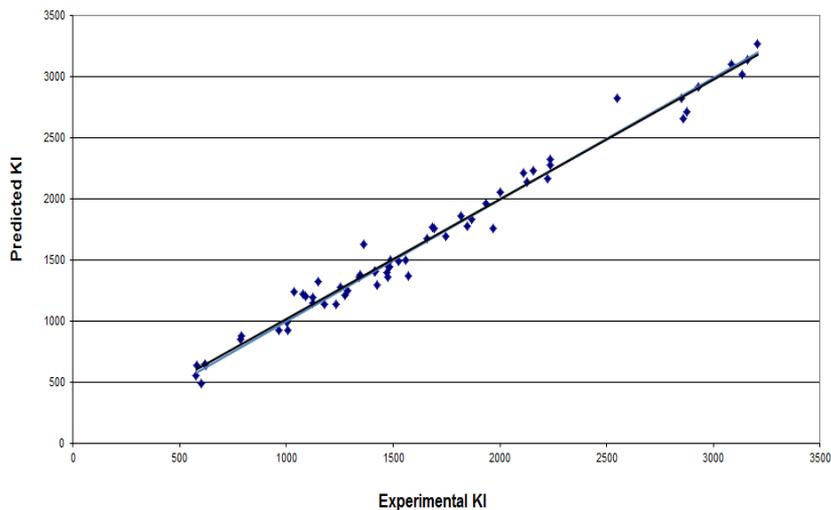
PMI RESEARCH & DEVELOPMENT

CASI Concept



Predictive QSPR Models for KI and 2DrelRT

Kovats Index

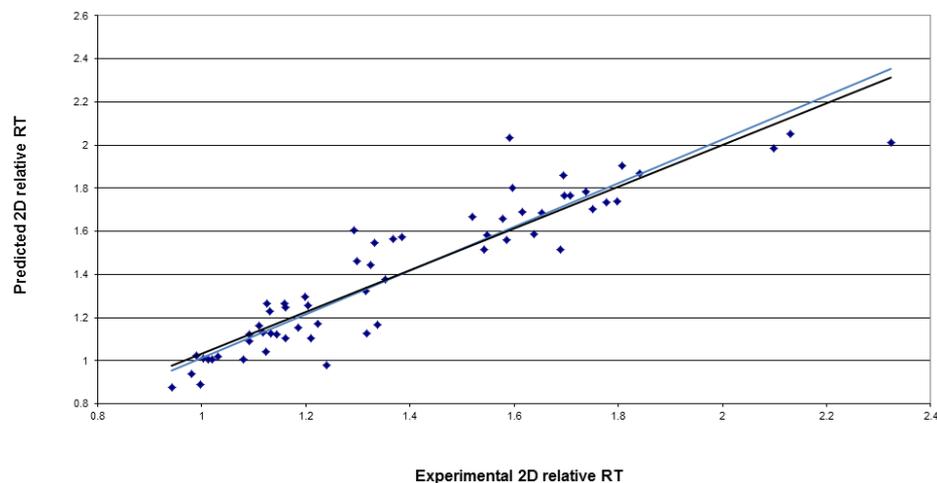


Validation n=60

$r^2 = 0.981$

GA – linear regression, 7 Molecular Descriptors

2DrelRT

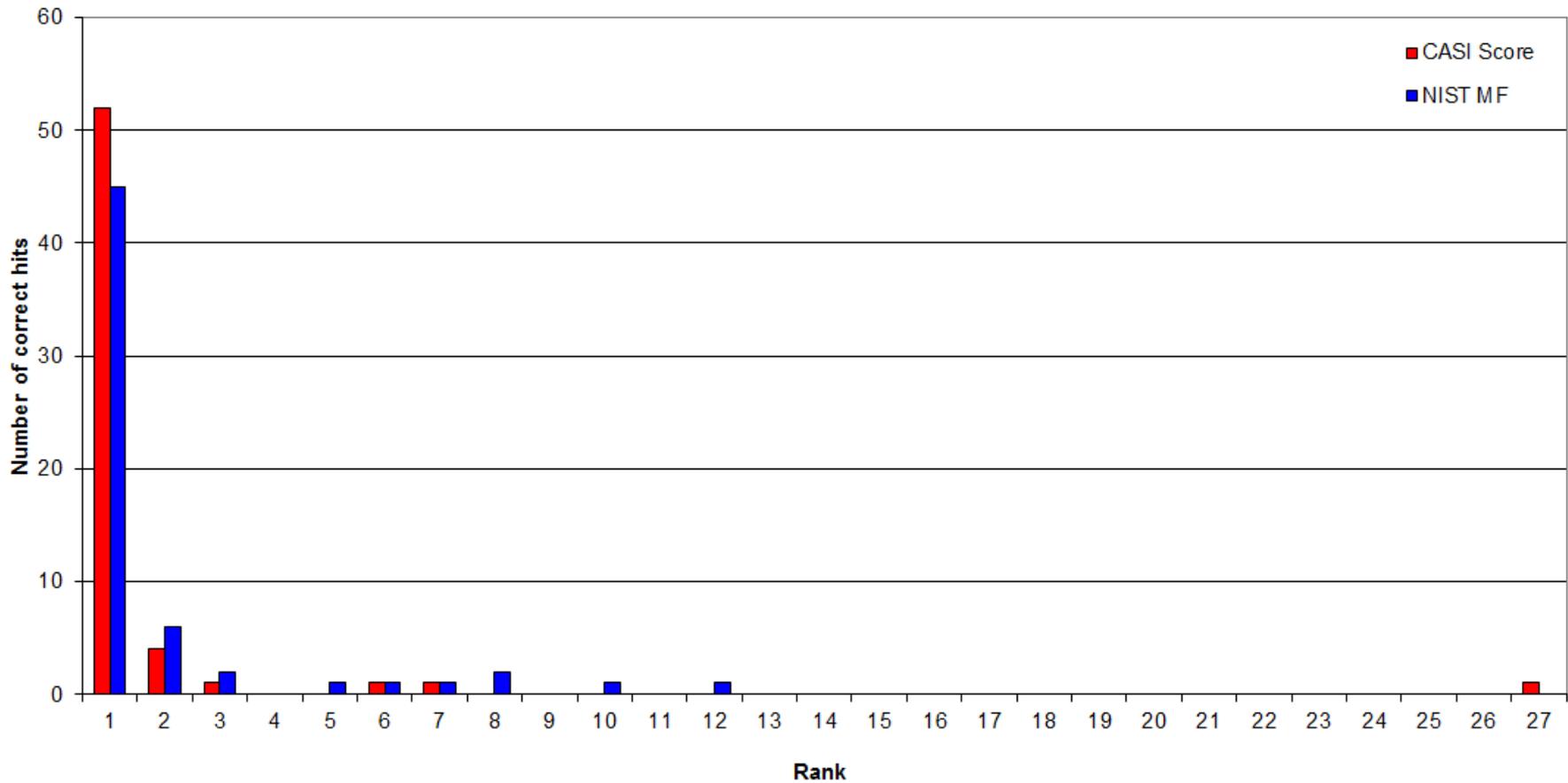


$r^2 = 0.855$

GA – Support Vector Regression, 12 molecular descriptors



Ranking of hits using CASI Score vs. NIST MS Match Factor



CASI Score improves NIST MS Search ranking.

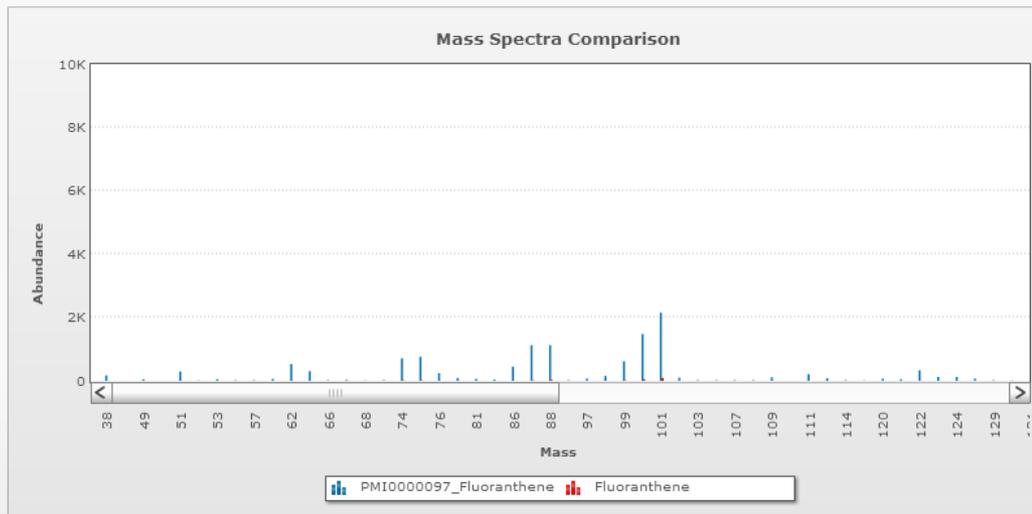


Spectra Comparison

Discriminatory Power: 1.01

Retention Time 1: 2129 Retention Time 2 (s): 1.84

BP (predicted): 361.51



Hit List

	PMI Code	CAS	Name	Match Factor	Iso. Mass	Score	Pred RT1	Pred RT2	List Marker	BP pred. (var)
	PMI0003510	206440	Fluoranthene	820	202.07825	819	2135.84 (.3 %)	1.87 (1.6 %)	-	375 (3.7 %)
	PMI0003509	129000	Pyrene	819	202.07825	809	2144.67 (.7 %)	1.91 (4 %)	-	404 (11.8 %)



CIKB Scope

Central platform to **report** on the presence and concentration of constituents in smoke aerosols and their known or measured toxicities to support the toxicological assessment.

Information of interest to be combined with chemical – analytical information of UCSD and CASI platforms:



Product Lifecycle: Complete specification of developed and tested products.



LIMS: Quantitative smoke characterization.



***In-vitro* Assays:** Standardized *in vitro* assays to assess toxicity.



Toxicities: Known toxicological information from literature.



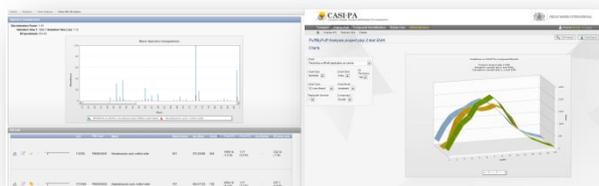
Metabolites: Known metabolites from literature.



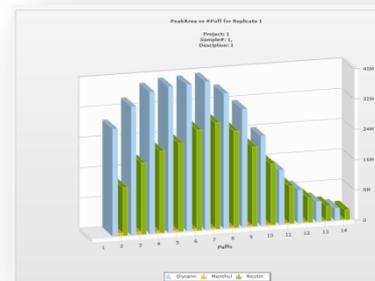
Chemistry Based Reporting



1. Unique Compounds & Spectra Database



2. Computer Assisted Structure Identification



3. Chemoinformatics Knowledge Base



Scientific Data Warehouse



Product Lifecycle



LIMS



Assays



Toxicities

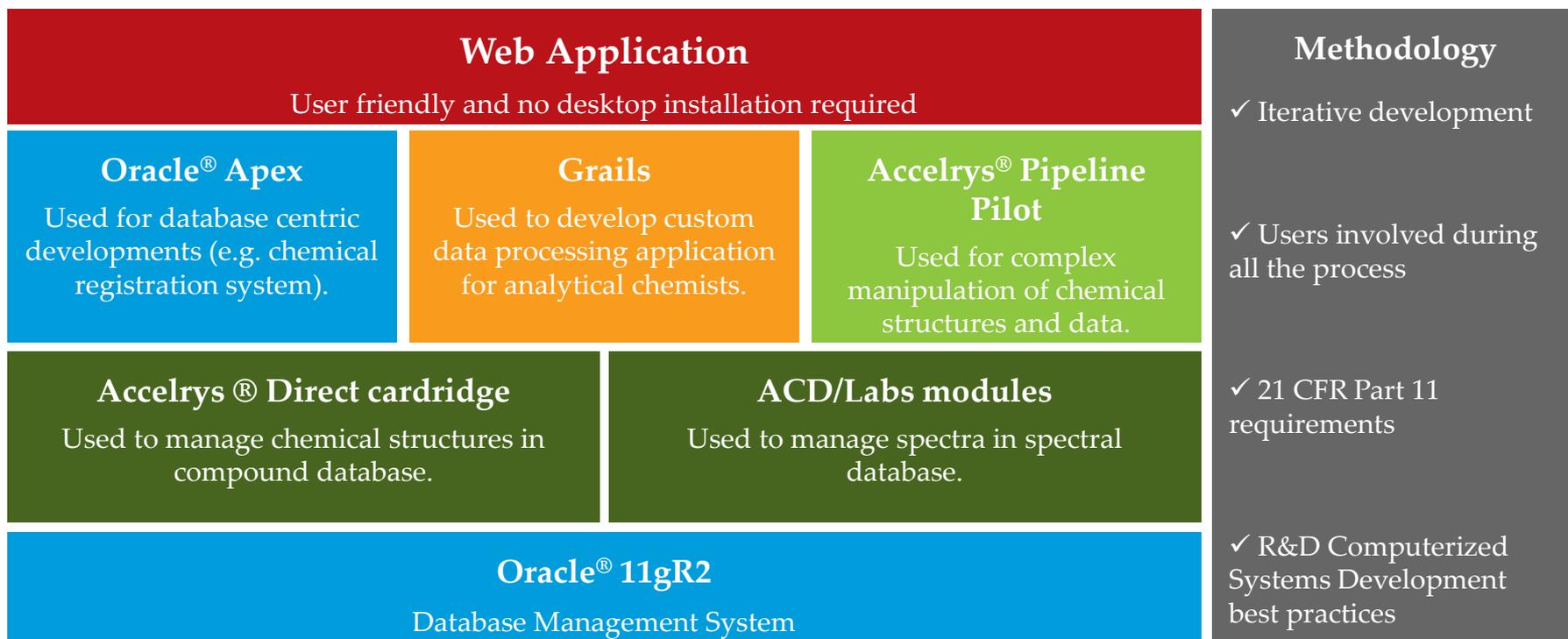


Metabolites



Used tools

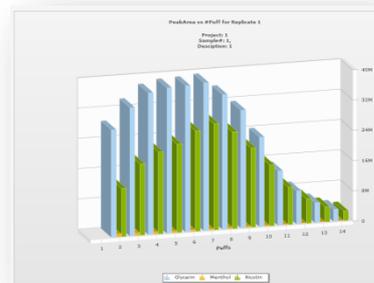
- Rapid Application Development tools were used to develop software to support the processing of analytical chemistry data (mass spectrometry) and thus to answer quickly to business needs.



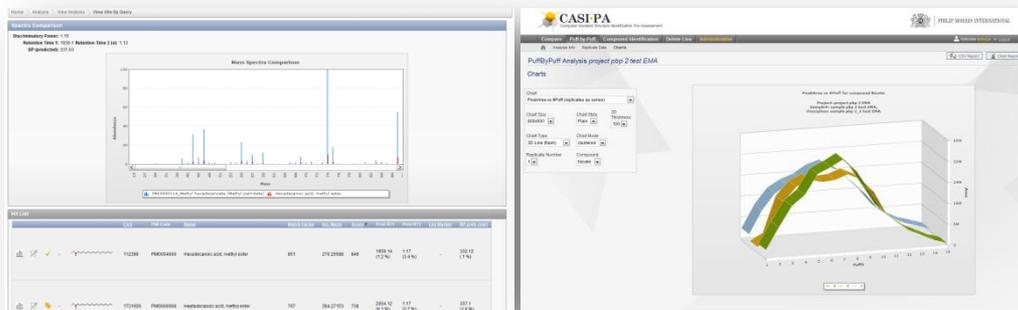
Conclusion



UCSD developed to facilitate and automatize processes of the unambiguous registration of compounds and their analytical spectra



CIKB in development to report on product chemical constituents and the associated toxicity and metabolic data



CASI platforms developed to increase the confidence in the identification of compounds using GC-MS methods and facilitates to users the comparison of analyses



Scientific Data Warehouse



Acknowledgment

- *Stephane Cano, Stephane Albrecht, Pavel Pospisil*
- *PMI R&D chemistry analytics teams*
- *PMI R&D HPC team*
- *PMI R&D SIS team*
- *Accelrys Symyx consultants*
- *ACD/Labs consultants*

Thank you for your attention.

