# MUNI RECETOX

# E2020 – Soft skills II – Information Literacy 3. Databases

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### I want to start working/writing, but ...

- What resources to use?
- Where can I find them?
- How can I find what interest me?
- What to do if I don't find what I need?
- How to formulate a query understood by the database?

### **Content**

- Lecture divided into 2 parts:
  - 1<sup>st</sup> part: Theoretical to understand how is the information stored

how to access it

- 2<sup>nd</sup> part: Practical – you are at the helm and you find the information!

### **VPN**

- MUNI VPN
  - MUNI
  - Eduroam

https://it.muni.cz/en/services/wireless-wi-fi-connection

#### Wireless Wi-Fi Connection

A majority of Masaryk University's premises is covered by Wi-Fi network enabling internet connection from laptops, tablets, and mobile phones. MU uses *Eduroam*, a world-wide network, which enables internet connection at most of academic institutions around the world based on the principle of unified login.

# #1 Eduroam (Main Network)

△ UČO¹)@muni.cz

secondary password<sup>2)</sup>

Eduroam is the main wireless network at MU enabling connection to internet. This network is being used by a variety of academic institutions around the world, so you can often connect to it automatically even on study or business trips.

# **#2 MUNI** (Auxiliary Network)

≗ UČO¹)

secondary password<sup>2)</sup>

Be careful, data transfer via this network is not secured! MUNI network functions as an auxiliary network. You should only use it if having trouble with Eduroam network.

### **Database**

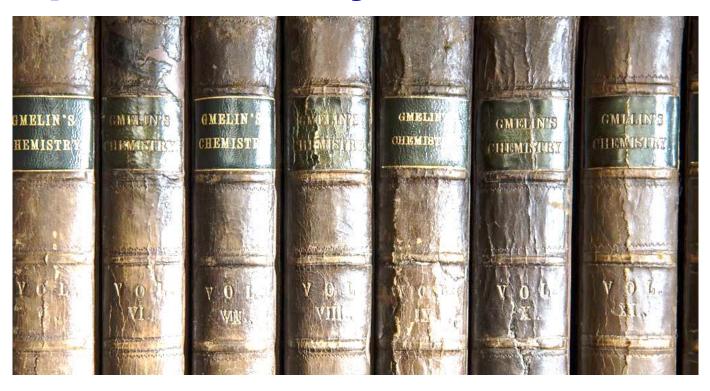
"a usually large collection of data organized especially for rapid search and retrieval (as by a computer)" (Webster dictionary)

Exist for multiples types of information

- Scientific Articles and Journals (WoS Scopus as seen in 2.Scientometry)
- Chemical and other information
- Specific properties and parameters

# Database: From "paper" to today

- First chemical database
  - First edition published in 1817,
     last edition: 8th edition 1990's



Nowadays most of them are digitalized and the information is available online

### **Chemical databases**

- Multiple existing databases based on different criterias
  - according to chemical structures
  - literature
  - crystallographic
  - spectroscopic: infrared, absorbance, nuclear magnetic resonance, ...
  - reactions
  - thermodynamic
  - others, …

### **Chemical databases**

- Often based on chemical structures searches
- Chemical structures are easy to read by humans (visual reading, ...)

1,5-Dihydroxy-4,8-dinitro-9,10-dioxo-9,10-dihydroanthracene-2,6-disulfonic acid

— How to explain it to computer / software / app ?

### Chemical structures and their representation

In order to look for a structure and for the computer to understand



- b) Connectivity matrices (e.g.MDL molfile, PDB, CML, ... = Computer languages related to chemistry created specifically for computation
- c) Linear strings: e.g. SMILES/SMARTS, SLN, WLN, InChi = easy computer language which computer can decipher the structure

### Chemical name (or similar unique identifier)

- Each compound can have multiple names
  - Common name (trivial, semi-trivial, systematic, business, ...)
    - Chlorpyrifos (Insecticide, banned since 2020 in Europe)
  - Proper chemical name: (= IUPAC name, International Union of Pure and Applied Chemistry)
    - -0,0-diethyl 0-(3,5,6-trichloro-2-pyridinyl)-phosphorothioate
  - Other names (sometimes names of commercial products they are featured in)
    - Chlorpyrifos-ethyl, Brodan, Bolton insecticide, Cobalt, ...

### Chemical name (or similar unique identifier)

- In order to remove possible errors and mistakes:
  - A unique numerical identifier was created =
  - <u>CAS RN</u> (Chemical Abstracts Service Registry Number)
  - Assigned to every chemical substance
  - From 1800's to today, registry account from more than 193 million compounds

### Chemical name (or similar unique identifier)

– Chlorpyrifos: CAS RN = 2921-88-2

#### Metabolites and derivatives:

– Chlorpyrifos-methyl = 5598-13-0

— Chlorpyrifos-oxon: 5598-15-2

### **Connectivity Matrix**

#### Computer language

- Chemical Table File (CT File)
  - Family of text based chemical file formats that describe molecules and chemical reactions
  - Numerous file format exist

- CT File is an open format
  - Lists each atom in a molecule, with the x-y-z coordinates of that atom, and the bonds amongst atoms
  - Just need to register on this website to access them: <a href="https://discover.3ds.com/ctfile-">https://discover.3ds.com/ctfile-</a> documentation-request-form

**Connectivity Matrix** 

Chiral

#### Molfile:

An MDL Molfile is a file format

- Contains information about:
  - atoms,
  - bonds, (=connectivity)
  - charges
  - coordinates of a molecule
- Recognized by most cheminformatics software systems/applications

L-Alanine	Title line (can be blank but line must exist)	
ABCDEFGH09071717443D	Program / file timestamp line (Name of source program and a file timestamp)	Header Block (3 lines)
Exported	Comment line (can be blank but line must exist)	
6 5 0 0 1 0 3 V2000	Counts line	
-0.6622 0.5342 0.0000 C 0 0 2 0 0 0 0.6622 -0.3000 0.0000 C 0 0 0 0 0 0 0 0 0 0 0 0 0 0	Atom block (1 line for each atom): x, y, z (in angstroms), element, etc.	Connection table
1 2 1 0 0 0 1 3 1 1 0 0 1 4 1 0 0 0 2 5 2 0 0 0 2 6 1 0 0 0	Bond block (1 line for each bond): 1st atom, 2nd atom, type, etc.	
M CHG 2 4 1 6 -1 M ISO 1 3 13	Properties block	
M END	END line (NOTE: some programs don't like a blank line before M END)	END

### **Connectivity Matrix**

#### Same exist for Proteins = PDB format

```
HEADER
         EXTRACELLULAR MATRIX
                                               22-JAN-98 1A3I
TITLE
         X-RAY CRYSTALLOGRAPHIC DETERMINATION OF A COLLAGEN-LIKE
        2 PEPTIDE WITH THE REPEATING SEQUENCE (PRO-PRO-GLY)
TITLE
EXPDTA
         X-RAY DIFFRACTION
AUTHOR
         R.Z.KRAMER, L.VITAGLIANO, J.BELLA, R.BERISIO, L.MAZZARELLA,
        2 B.BRODSKY, A.ZAGARI, H.M.BERMAN
AUTHOR
REMARK 350 BIOMOLECULE: 1
REMARK 350 APPLY THE FOLLOWING TO CHAINS: A, B, C
REMARK 350
            BIOMT1 1 1.000000 0.000000 0.000000
                                                          0.00000
           BIOMT2 1 0.000000 1.000000 0.000000
REMARK 350
                                                          0.00000
SEQRES
       1 A
              9 PRO PRO GLY PRO PRO GLY PRO PRO GLY
       1 B
               6 PRO PRO GLY PRO PRO GLY
SEQRES
       1 C
               6 PRO PRO GLY PRO PRO GLY
                               8.316 21.206 21.530 1.00 17.44
ATOM
                PRO A 1
MOTA
         2 CA PRO A 1
                               7.608 20.729 20.336 1.00 17.44
                PRO A 1
                               8.487 20.707 19.092 1.00 17.44
ATOM
                PRO A 1
                               9.466 21.457 19.005 1.00 17.44
ATOM
ATOM
         5 CB
                PRO A 1
                               6.460 21.723 20.211 1.00 22.26
                               3.682 22.541 11.236 1.00 21.19
                                                                         C
HETATM 130 C
      131 0
                               2.807 23.097 10.553 1.00 21.19
HETATM 132 OXT ACY
                               4.306 23.101 12.291 1.00 21.19
```

## Connectivity Matrix without connectivity

#### X-Y-Z file

- No information about bonds (covalent, hydrogens, VdW, ...) admits a greater flexibility
- Typical XYZ format specifies the molecule geometry
  - First line = number of atoms with Cartesian coordinates
  - Second line = a comment
  - Third and following line = atomic coordinates

```
<number of atoms>
comment line
atom_symbol<sub>11</sub> x-coord<sub>11</sub> y-coord<sub>11</sub> z-coord<sub>11</sub>
atom_symbol<sub>12</sub> x-coord<sub>12</sub> y-coord<sub>12</sub> z-coord<sub>12</sub>
atom_symbol<sub>in</sub> x-coord<sub>in</sub> y-coord<sub>in</sub> z-coord<sub>in</sub>
<number of atoms>
comment line
atom_symbol<sub>21</sub> x-coord<sub>21</sub> y-coord<sub>21</sub> z-coord<sub>21</sub>
atom_symbol<sub>22</sub> x-coord<sub>22</sub> y-coord<sub>22</sub> z-coord<sub>22</sub>
```

### **Connectivity Matrix without connectivity**

#### Pyridine

Formula: C<sub>5</sub>H<sub>5</sub>N

$$H \setminus_{C \setminus_{C}} H$$
 $H \setminus_{C \setminus_{C}} H$ 
 $H \setminus_{C \setminus_{C}} H$ 

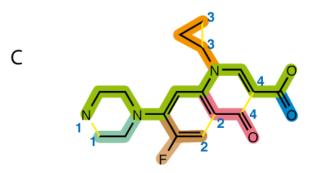
C C	-0.180226841 -0.180226841 -0.180226841 -0.180226841	0.360945118 1.559292118 1.503191118 0.360945118	-1.120304970 -0.407860970 0.986935030 1.29018350
С	-0.180226841	1.503191118	0.986935030
_			
N	-0.180226841	0.360945118	1.29018350
14			
С	-0.180226841	-0.781300882	0.986935030
С	-0.180226841	-0.837401882	-0.407860970
Н	-0.180226841	0.360945118	-2.206546970
Н	-0.180226841	2.517950118	-0.917077970
Н	-0.180226841	2.421289118	1.572099030
Н	-0.180226841	-1.699398882	1.572099030
Н	-0.180226841	-1.796059882	-0.917077970

**Linear string**: represents structures as a linear string of characters

#### Simplified Molecular Input Line Entry Specification (SMILES)

- Chemical notation allowing user to represent a chemical structure
- Easily read, understood and used by computer
- Contains connectivity, but no longer 2D or 3D coordinates

- How does it work?
  - Every atoms are supported
  - Upper-case for aromatic atoms, lowercase for non-aromatic atoms
  - Bonds:
    - single bond
    - = double bond
    - # triple bond
    - \* aromatic bond
    - . disconnected structures



D

N1CCN(CC1)C(C(F)=C2)=CC(=C2C4=O)N(C3CC3)C=C4C(=O)O

#### Simple chain molecule

(Hydrogen suppressed = no need to put hydrogen in it, software understand that they are here)

SMILES	Formula	Name	Structure
CC	CH <sub>3</sub> CH <sub>3</sub>	Ethane	H H H – C — H H – H
C=C	CH <sub>2</sub> CH <sub>2</sub>	Ethene	H H C=C H H
	CH <sub>3</sub> Br	Bromomethane	H H H

#### Simple chain molecule

(Hydrogen suppressed = no need to put hydrogen in it, software understand that they are here)

SMILES	Formula	Name	Structure
CC	CH <sub>3</sub> CH <sub>3</sub>	Ethane	H H I I H – C —— C – H I I H H
C=C	CH <sub>2</sub> CH <sub>2</sub>	Ethene	H H C=C H H
CBr	CH₃Br	Bromomethane	H H H

#### **Branches**

(in parentheses = a branche placed right after the atom it is connected to)

SMILES	Formula	Name	Structure
CC(O)C	CH <sub>3</sub> CHOHCH <sub>3</sub>	2-propanol	н он н Н О О Н Н О О Н Н Н Н Н
CC(=O)C	CH₃COCH₃	2-propanone (acetone)	H O H 
	CH <sub>3</sub> CH <sub>3</sub> CHCH <sub>2</sub> CH <sub>3</sub>	2-methylbutane (Isopentane)	CH <sub>3</sub> H 

#### **Branches**

(in parentheses = a branche placed right after the atom it is connected to)

SMILES	Formula	Name	Structure
CC(O)C	CH <sub>3</sub> CHOHCH <sub>3</sub>	2-propanol	H H H H H H H H H H H H H H H H H H H
CC(=O)C	CH <sub>3</sub> COCH <sub>3</sub>	2-propanone (acetone)	H-C-H H-C-H H-C-H
CCC(C)C	CH <sub>3</sub> CH <sub>3</sub> CHCH <sub>2</sub> CH <sub>3</sub>	2-methylbutane (Isopentane=	CH <sub>3</sub> H

#### Rings:

(Use number to identify opening and closing of ring atom)

SMILES	Formula	Name	Structure
C=1CCCCC1 Also C*1*C*C*C*C*C1	CHCHCH <sub>2</sub> CH <sub>2</sub> CH <sub>2</sub> CH	Cyclohexene	
	CH <sub>2</sub> (O)CHCH <sub>2</sub> CH <sub>3</sub>	Ethyloxirane	CH <sub>3</sub>
	СНСНСНСНСНСНСНСНСН	Naphtalene	1 2

#### Rings:

(Use number to identify opening and closing of ring atom)

SMILES	Formula	Name	Structure
C=1CCCCC1 Also C*1*C*C*C*C*C1	CHCHCH <sub>2</sub> CH <sub>2</sub> CH <sub>2</sub> CH	Cyclohexene	
C1OC1CC	CH <sub>2</sub> (O)CHCH <sub>2</sub> CH <sub>3</sub>	Ethyloxirane	CH <sub>3</sub>
c1cc2cccc2cc1	СНСНСНСНСНСНСНСНСН	Naphtalene	1 2

#### **Charged atoms:**

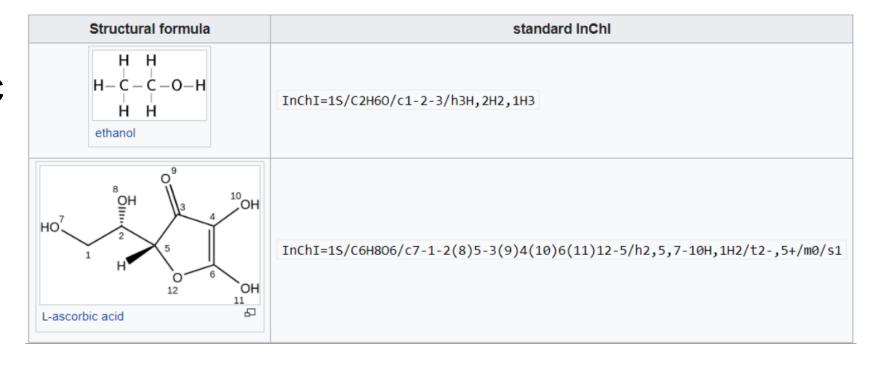
(Atoms followed by brackets which enclose the charge on the atom, maybe be explicitly stated ({-1}) or not ({-}))

SMILES	Name	Structure
CCC(=O)O{-1}		
Or	lonised form of propanoic acid	
CCC(=O)O{-}		0
c1ccccn{+1}1CC(=O)O	1-Carboxylmethyl pyridinium	N <sup>+</sup> OH

## **Linear string: InChl**

#### <u>InChl</u>

- InternationalChemical Identifier
- Introduced by IUPACas a **standard** in2006
- Contains different layers:
  - general formula,
  - hydrogens, charges,
  - stereochemistry,
  - isotopes,...



### Representation of structure - Conclusions

- Chemical names connectivity matrices linear strings
- Multiple choice exist, none of them wrong, some more popular than others

File Extension	MIME Type	Proper Name	Description
alc	chemical/x-alchemy	Alchemy Format	
csf	chemical/x-cache-csf	CAChe MolStruct CSF	
cbin, cascii, ctab	chemical/x-cactvs-binary	CACTVS format	
cdx	chemical/x-cdx	ChemDraw eXchange file	
cer	chemical/x-cerius	MSI Cerius II format	
c3d	chemical/x-chem3d	Chem3D Format	
chm	chemical/x-chemdraw	ChemDraw file	
cif	chemical/x-cif	Crystallographic Information File, Crystallographic Information Framework	Promulgated by the International Union of Crystallography
cmdf	chemical/x-cmdf	CrystalMaker Data format	
cml	chemical/x-cml	Chemical Markup Language	XML based Chemical Markup Language.
сра	chemical/x-compass	Compass program of the Takahashi	
bsd	chemical/x-crossfire	Crossfire file	
csm, csml	chemical/x-csml	Chemical Style Markup Language	
ctx	chemical/x-ctx	Gasteiger group CTX file format	
cxf, cef	chemical/x-cxf	Chemical eXchange Format	
emb, embl	chemical/x-embl-dl-nucleotide	EMBL Nucleotide Format	
spc	chemical/x-galactic-spc	SPC format for spectral and chromatographic data	

### **Questions?**

- Contact me anytime via email: <a href="mailto:ludovic.mayer@recetox.muni.cz">ludovic.mayer@recetox.muni.cz</a>