

Molecular Similarity and Chemical Similarity



SAPIENZA
UNIVERSITÀ DI ROMA



Overview

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Chemical and Molecular Similarity

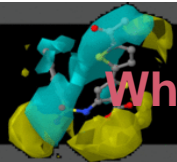
Molecular similarity importance

Representations of chemical structures

Similarity and Dissimilarity measures

Similarity searching

Cluster and diversity analysis



What does similarity stand for?

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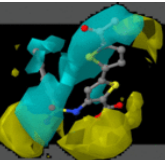


The terms chemical and molecular similarity are often used synonymously but this may not be entirely accurate.

Chemical similarity is based primarily on the physicochemical characteristics of compounds (e.g., solubility, boiling point, log P, molecular weight, electron densities, dipole moments, etc.)

Molecular similarity focuses primarily on the structural features (e.g., shared substructures, ring systems, topologies, etc.) of compounds and their representation.

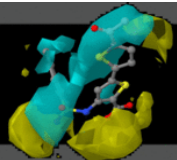
Similarity perception



Chemical similarity	Mol. weight	LogP	Rotatable bonds	Aromatic rings	Heavy atoms	
	A	341.4	5.23	4	4	26
	B	463.5	4.43	4	5	35
Molecular similarity						
2D similarity						
3D similarity						

Two *Vascular Endothelial Growth Factor Receptor 2* ligands and different ways to assess their similarity.

Biological similarity	Vascular endothelial growth factor receptor 2	Tyrosine-protein kinase TIE-2	
	A	active	inactive
	B	active	active
Global similarity			
Local similarity			



The notion of chemical similarity (or molecular similarity) is one of the most important concepts in chemoinformatics.

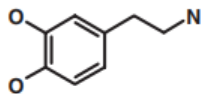
It plays an important role in predicting the properties of chemical compounds, designing chemicals with a predefined set of properties and, especially, in conducting drug design studies by screening large databases containing structures of available (or potentially available) chemicals.

There are many ways to measure the similarity

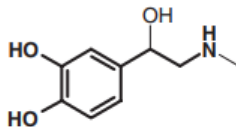


Substructure search

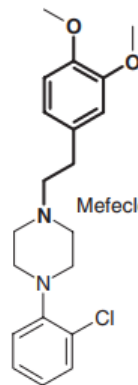
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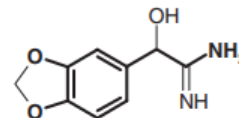
query



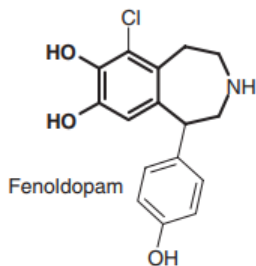
Adrenaline



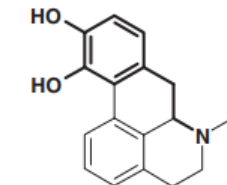
Mefeclozazine



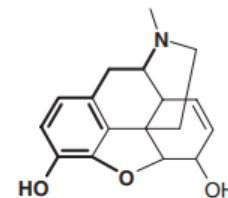
Olmidine



Fenoldopam



Apomorphine



Morphine



Neighbourhood principle

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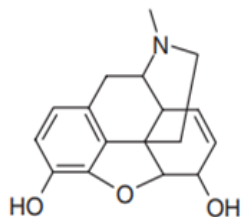
The rationale for similarity searching lies in the similar property principle [Johnson and Maggiora 1990] which states that structurally similar molecules tend to have similar properties.

Given a molecule of known biological activity, compounds that are structurally similar to it are likely to exhibit the same activity. This characteristic has been referred to as neighbourhood behaviour [Patterson et al. 1996].

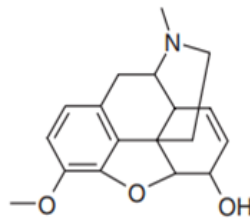


Neighbourhood principle (2)

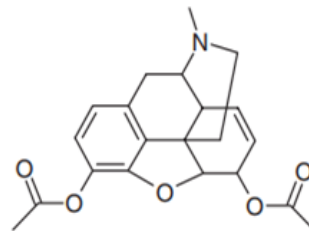
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Morphine

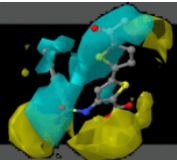


Codeine



Heroin

Compounds active at opioid receptors

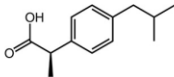
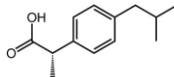
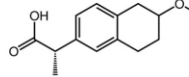
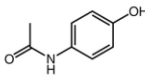
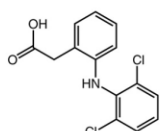
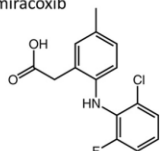


Neighbourhood principle (3)

Compounds such as the ibuprofen enantiomers, ibuprofen and paracetamol, or diclofenac and lumiracoxib, appear visibly similar.

From a medicinal chemistry point of view, however, this assessment may not be generally agreed upon since small chemical differences can lead to important changes in specificity profiles (e.g., diclofenac vs lumiracoxib) or compounds containing different functional groups can be synthesized or derivatized in different ways (e.g., ibuprofen vs paracetamol).

Moreover, these COX inhibitors are involved in highly complex similarity–activity relationships (similar mechanism of action) that also cannot easily be separated from a medicinal chemistry perspective.

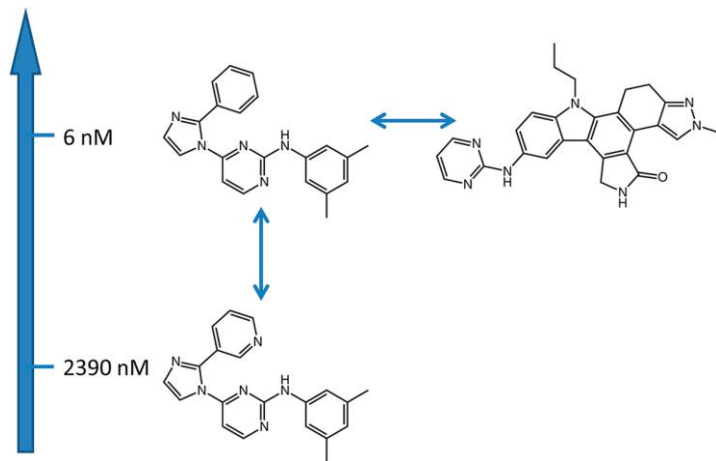
<p>(R)-(-)-Ibuprofen</p>  <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>no</td> <td>no</td> <td>no</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	no	no	no	<p>(S)-(+)-Ibuprofen</p>  <p>Bioavailability 49 -73%</p> <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>yes</td> <td>yes</td> <td>no</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	yes	yes	no	<p>(S)-(+)-Naproxen</p>  <p>Bioavailability 95%</p> <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>yes</td> <td>yes</td> <td>yes</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	yes	yes	yes
COX-1	COX-2	HSL																		
no	no	no																		
COX-1	COX-2	HSL																		
yes	yes	no																		
COX-1	COX-2	HSL																		
yes	yes	yes																		
<p>Paracetamol</p>  <p>Bioavailability ~100%</p> <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>no</td> <td>yes</td> <td>no</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	no	yes	no	<p>Diclofenac</p>  <p>Bioavailability >99%</p> <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>yes</td> <td>yes</td> <td>no</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	yes	yes	no	<p>Lumiracoxib</p>  <p>Bioavailability ~74%</p> <table border="1"> <thead> <tr> <th>COX-1</th> <th>COX-2</th> <th>HSL</th> </tr> </thead> <tbody> <tr> <td>no</td> <td>yes</td> <td>no</td> </tr> </tbody> </table>	COX-1	COX-2	HSL	no	yes	no
COX-1	COX-2	HSL																		
no	yes	no																		
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COX-1	COX-2	HSL																		
no	yes	no																		

Cyclooxygenase (COX) inhibitors and their activity profiles are compared.
HSL = Hormone-Sensitive Lipase



Neighbourhood principle exceptions

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Similarity versus activity. Three vascular endothelial growth factor receptor 2 ligands are shown that represent different (vertical vs horizontal) similarity-activity (potency) relationships.

There are many exceptions to the principle but it is an excellent rule-of-thumb in the absence of more detailed knowledge

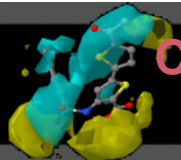
People's judgements of similarity are inherently subjective, so need to provide a quantitative basis, a similarity measure, for assessing the degree of resemblance



The main difficulty with similarity searching is that assessing the degree of similarity between two objects is subjective.

In order to be able to quantify the similarity between two molecules, a similarity searching method requires two components:

- **a set of numerical descriptors** that can be used to compare molecules;
- **a similarity coefficient** which provides a way of quantifying the degree of similarity based on the descriptors.



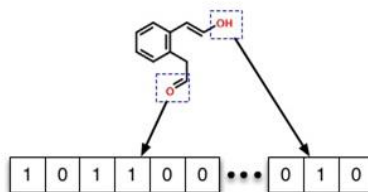
Numerical values assigned to structures:

- 1D properties: MW, logP, PSA etc
- 2D properties: fingerprints, topological indices
- Maximum Common Substructures
- 3D properties: molecular fields, shape



Molecular fingerprints

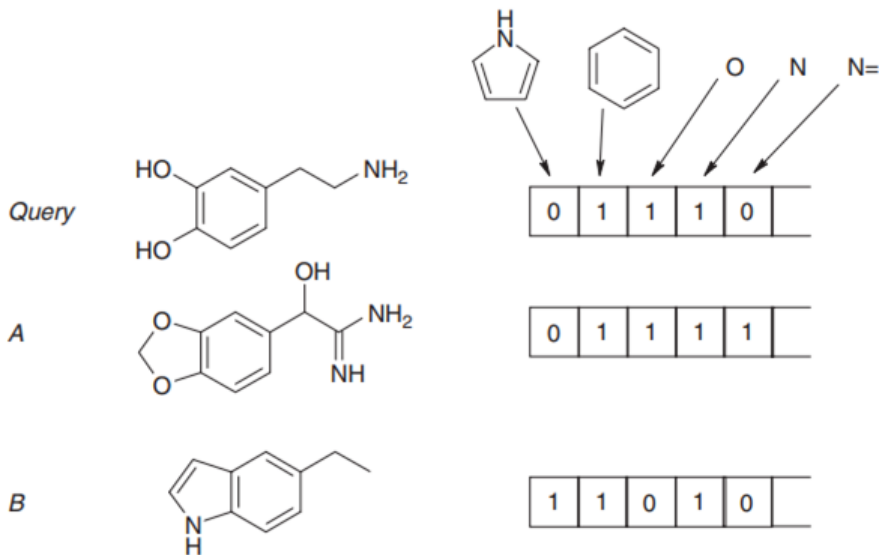
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- Lots of types of fingerprints
- “Keyed” fingerprints indicate the presence or absence of a structural feature
- Length can vary from 166 to 4096 bits or more
- Fingerprints usually compared using the Tanimoto metric



Fingerprints: an example



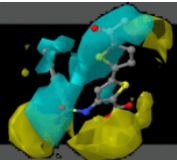


Similarity measures

Name	Formula for continuous variables	Formula for binary (dichotomous) variables
Tanimoto (Jaccard) coefficient	$S_{AB} = \frac{\sum_{i=1}^N x_{iA} x_{iB}}{\sum_{i=1}^N (x_{iA})^2 + \sum_{i=1}^N (x_{iB})^2 - \sum_{i=1}^N x_{iA} x_{iB}}$ Range: -0.333 to +1	$S_{AB} = \frac{c}{a+b-c}$ Range: 0 to 1
Dice coefficient (Hodgkin index)	$S_{AB} = \frac{2 \sum_{i=1}^N x_{iA} x_{iB}}{\sum_{i=1}^N (x_{iA})^2 + \sum_{i=1}^N (x_{iB})^2}$ Range: -1 to +1	$S_{AB} = \frac{2c}{a+b}$ Range: 0 to 1
Cosine similarity (Carbó index)	$S_{AB} = \frac{\sum_{i=1}^N x_{iA} x_{iB}}{[\sum_{i=1}^N (x_{iA})^2 \sum_{i=1}^N (x_{iB})^2]^{1/2}}$ Range: -1 to +1	$S_{AB} = \frac{c}{\sqrt{ab}}$ Range: 0 to 1
Euclidean distance	$D_{AB} = [\sum_{i=1}^N (x_{iA} - x_{iB})^2]^{1/2}$ Range: 0 to ∞	$D_{AB} = \sqrt{a+b-2c}$ Range: 0 to N
Hamming (Manhattan or City-block) distance	$D_{AB} = \sum_{i=1}^N x_{iA} - x_{iB} $ Range: 0 to ∞	$D_{AB} = a + b - 2c$ Range: 0 to N
Soergel distance	$D_{AB} = \frac{\sum_{i=1}^N x_{iA} - x_{iB} }{\sum_{i=1}^N \max(x_{iA}, x_{iB})}$ Range: 0 to 1	$D_{AB} = \frac{a+b-2c}{a+b-c}$ Range: 0 to 1

- Similarity (**S**) or distance (**D**) coefficients in common use for similarity searching in chemical databases.
- For binary data:
 - **a** is defined as the number of bits set to "1" in molecule **A**
 - **b** as the number of bits set to "1" in molecule **B**
 - **c** as the number of bits that are "1" in both **A** and **B**.

[Willett et al. 1998.]



Basic bit count terms of similarity

Symbol	Description																	
$onlyA $	number of bits set "on" in fingerprint A but not in B	A <table border="1"><tr><td>1</td><td>0</td><td>1</td><td>1</td><td>1</td><td>0</td><td>0</td><td>0</td></tr></table> B <table border="1"><tr><td>1</td><td>1</td><td>0</td><td>1</td><td>1</td><td>0</td><td>1</td><td>0</td></tr></table>	1	0	1	1	1	0	0	0	1	1	0	1	1	0	1	0
1	0	1	1	1	0	0	0											
1	1	0	1	1	0	1	0											
$onlyB $	number of bits set "on" in fingerprint B but not in A	A <table border="1"><tr><td>1</td><td>0</td><td>1</td><td>1</td><td>1</td><td>0</td><td>0</td><td>0</td></tr></table> B <table border="1"><tr><td>1</td><td>1</td><td>0</td><td>1</td><td>1</td><td>0</td><td>1</td><td>0</td></tr></table>	1	0	1	1	1	0	0	0	1	1	0	1	1	0	1	0
1	0	1	1	1	0	0	0											
1	1	0	1	1	0	1	0											
$bothAB $	number of bits set "on" in both fingerprints	A <table border="1"><tr><td>1</td><td>0</td><td>1</td><td>1</td><td>1</td><td>0</td><td>0</td><td>0</td></tr></table> B <table border="1"><tr><td>1</td><td>1</td><td>0</td><td>1</td><td>1</td><td>0</td><td>1</td><td>0</td></tr></table>	1	0	1	1	1	0	0	0	1	1	0	1	1	0	1	0
1	0	1	1	1	0	0	0											
1	1	0	1	1	0	1	0											
$neitherAB $	number of bits set "off" in both fingerprints	A <table border="1"><tr><td>1</td><td>0</td><td>1</td><td>1</td><td>1</td><td>0</td><td>0</td><td>0</td></tr></table> B <table border="1"><tr><td>1</td><td>1</td><td>0</td><td>1</td><td>1</td><td>0</td><td>1</td><td>0</td></tr></table>	1	0	1	1	1	0	0	0	1	1	0	1	1	0	1	0
1	0	1	1	1	0	0	0											
1	1	0	1	1	0	1	0											
$ A $	number of bits set "on" in fingerprint A																	
$ B $	number of bits set "on" in fingerprint B																	



Euclidean

Formula:

$$Sim_{Euclid}(A, B) = \sqrt{\frac{bothAB+neitherAB}{onlyA+onlyB+bothAB+neitherAB}}$$

Range:

$$[0.0 - 1.0]$$

$$\begin{aligned} \mathbf{p} &= \mathbf{A} \\ \mathbf{q} &= \mathbf{B} \end{aligned}$$

$$d(\mathbf{p}, \mathbf{q}) = d(\mathbf{q}, \mathbf{p}) = \sqrt{(q_1 - p_1)^2 + (q_2 - p_2)^2 + \dots + (q_n - p_n)^2}$$

Example:

$$= \sqrt{\sum_{i=1}^n (q_i - p_i)^2}$$

A	1	0	1	1	1	0	0	0
B	1	1	0	1	1	0	1	0

$$\sqrt{\frac{bothAB+neitherAB}{onlyA+onlyB+bothAB+neitherAB}} = \sqrt{\frac{3+1}{1+2+3+1}} = \sqrt{\frac{4}{8}} = 0.707$$



Manhattan

Formula:

$$Sim_{Manhattan}(A, B) = \frac{onlyA+onlyB}{onlyA+onlyB+bothAB+neitherAB}$$

Range:

$$[1.0 - 0.0]$$

Example:

A	1	0	1	1	1	0	0	0
B	1	1	0	1	1	0	1	0

$$\frac{onlyA+onlyB}{onlyA+onlyB+bothAB+neitherAB} = \frac{1+2}{1+2+3+2} = \frac{3}{8} = 0.375$$



Cosine

Formula:

$$Sim_{Cosine}(A, B) = \frac{bothAB}{\sqrt{(onlyA+bothAB)*(onlyB+bothAB)}}$$

Range:

[0.0 – 1.0]

$$similarity = \cos(\theta) = \frac{\mathbf{A} \cdot \mathbf{B}}{\|\mathbf{A}\| \|\mathbf{B}\|} = \frac{\sum_{i=1}^n A_i B_i}{\sqrt{\sum_{i=1}^n A_i^2} \sqrt{\sum_{i=1}^n B_i^2}}$$

Example:

A	1	0	1	1	1	0	0	0
B	1	1	0	1	1	0	1	0

$$\frac{bothAB}{\sqrt{(onlyA+bothAB)*(onlyB+bothAB)}} = \frac{3}{\sqrt{(1+3)*(2+3)}} = \frac{3}{\sqrt{20}} = 0.67$$

Calculates the ratio of the bits in common to the geometric mean of the number of "on" bits in the two fingerprints.



Tanimoto (Jaccard)

Formula:

$$Sim_{Tanimoto}(A, B) = \frac{bothAB}{|A|+|B|-bothAB} = \frac{bothAB}{onlyA+onlyB+bothAB}$$

Range:

[0.0 – 1.0]

$$J(A, B) = \frac{|A \cap B|}{|A \cup B|} = \frac{|A \cap B|}{|A| + |B| - |A \cap B|}$$

Example:

A	1	0	1	1	1	0	0	0
B	1	1	0	1	1	0	1	0

$$\frac{bothAB}{onlyA+onlyB+bothAB} = \frac{3}{1+2+3} = \frac{3}{6} = 0.5$$



Tversky

Formula:

$$Sim_{Tversky}(A, B) = \frac{bothAB}{\alpha * onlyA + \beta * onlyB + bothAB} \quad S(X, Y) = \frac{|X \cap Y|}{|X \cap Y| + \alpha |X - Y| + \beta |Y - X|}$$

The Tversky similarity measure is asymmetric. Setting the parameters $\alpha = \beta = 1.0$ is identical to using the *Tanimoto* measure.

The factor α weights the contribution of the first 'reference' molecule. The larger α becomes, the more weight is put on the bit setting of the reference molecule.

Range:

Setting $\alpha = \beta = 1$ produces the *Tanimoto coefficient*.

Setting $\alpha = \beta = 0.5$ produces *Dice's coefficient*.

variable

If we consider **A** to be the prototype and **B** to be the variant, then α corresponds to the weight of the prototype and β corresponds to the weight of the variant.

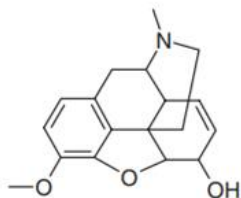
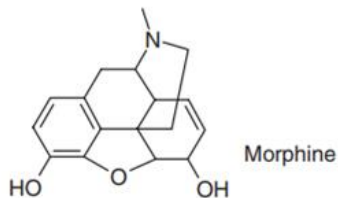
Example:

A	1	0	1	1	1	0	0	0
B	1	1	0	1	1	0	1	0

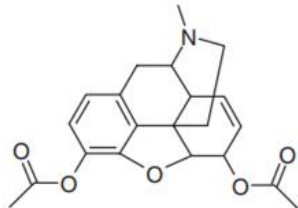
$$\frac{bothAB}{\alpha * onlyA + \beta * onlyB + bothAB} (\alpha = 2.0, \beta = 1.0) = \frac{3}{2.0 * 1 + 1.0 * 2 + 3} = \frac{3}{7} = 0.438$$



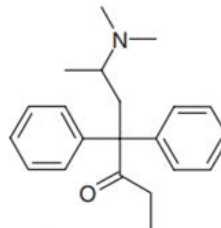
2D similarity (1)



Codeine
0.99 similar



Heroin
0.95 similar

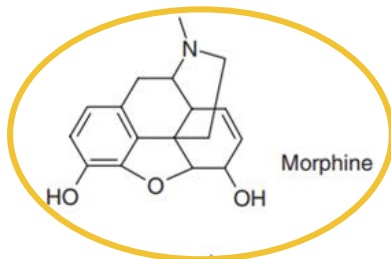


Methadone
0.20 similar

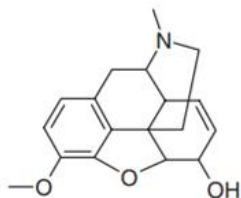
Similarities to morphine calculated using Daylight fingerprints and the Tanimoto coefficient.



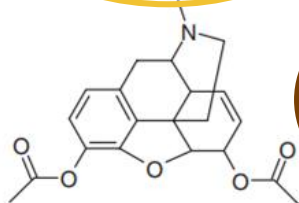
2D similarity (2)



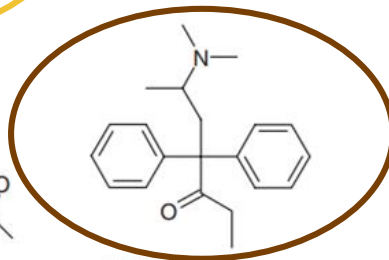
Molecular recognition depends on the 3D structure and properties (e.g. electrostatics and shape) of a molecule rather than the underlying substructure(s).



Codeine
0.99 similar



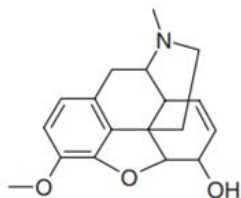
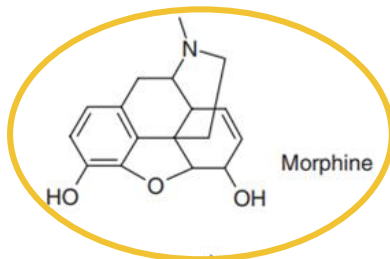
Heroin
0.95 similar



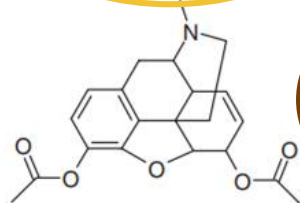
Methadone
0.20 similar



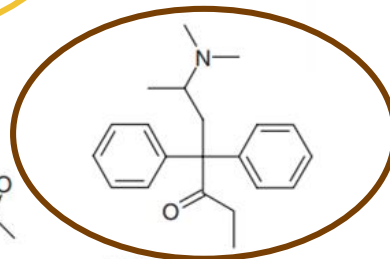
2D versus 3D Similarity



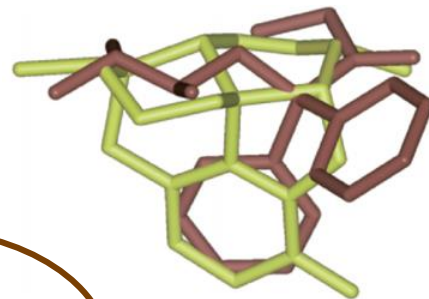
Codeine
0.99 similar



Heroin
0.95 similar



Methadone
0.20 similar

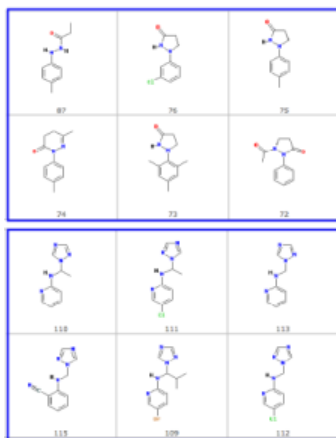




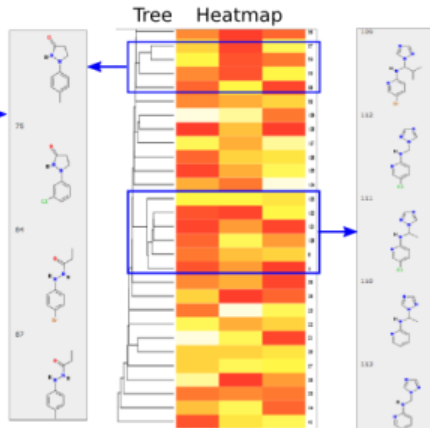
Cluster and diversity analysis

by **RCMD**.it

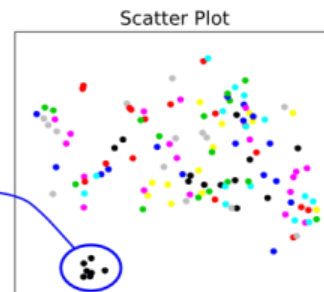
Binning Clustering

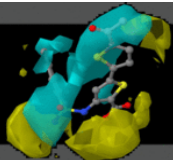


Hierarchical Clustering



Multidimensional Scaling





References

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- ❖ Baldi, Pierre, and Ramzi Nasr. "When is chemical similarity significant? The statistical distribution of chemical similarity scores and its extreme values." *Journal of chemical information and modeling* 50.7 (2010): 1205-1222.
- ❖ Riniker, Sereina, and Gregory A. Landrum. "Open-source platform to benchmark fingerprints for ligand-based virtual screening." *Journal of cheminformatics* 5.1 (2013): 1.
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- ❖ Duesbury, Edmund, John Holliday, and Peter Willett. "Maximum common substructure-based data fusion in similarity searching." *Journal of chemical information and modeling* 55.2 (2015): 222-230.
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- ❖ Schyman, Patric, Ruifeng Liu, and Anders Wallqvist. "General Purpose 2D and 3D Similarity Approach to Identify hERG Blockers." *Journal of chemical information and modeling* 56.1 (2016): 213-222.
- ❖ <http://www.daylight.com/dayhtml/doc/theory/theory.finger.html>
- ❖ <https://docs.eyesopen.com/toolkits/python/graphsimtk/measure.html>



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ChemMine Tools

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WORKBENCH

[My Compounds](#)

[Add Compounds](#)

TOOLS

[Past Jobs](#)

[Upload Numeric Data](#)

[Cluster](#)

[Physicochemical Properties](#)

[Similarity](#)

[Workbench](#)

Welcome!

To get started just add compounds.

(I) WORKBENCH

(II) SIMILARITY TOOLBOX

(IV) CLUSTERING TOOLBOX

(III) SEARCH TOOLBOX

EI SEARCH FINGERPRINT

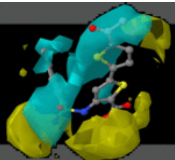
Pubchem

(V) PROPERTY TOOLBOX

Number	Weight	Charge	Number_of_Aromatic_Rings	Number_of_Aliphatic_Rings	Number_of_Cyclic_Rings	Number_of_Quaternary_Centers	Number_of_Tertiary_Centers	Number_of_Secondary_Centers	Number_of_Primary_Centers	Number_of_Hydrogen_Bond_Donors	Number_of_Hydrogen_Bond_Acceptors	Number_of_Rotatable_Bonds	Number_of_Cyclable_Bonds	Number_of_Cyclable_Centers
133	162.214	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
134	186.214	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
135	172.200	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
136	200.204	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
137	112.200	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
138	164.214	0.000	1.000	0.000	1.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000



The screenshot shows a web browser window with the address bar displaying "chemmine.ucr.edu/myCompounds/". The page has a dark header with the title "ChemMine Tools" and navigation links for "About", "Help", and "Downloads". Below the header, there is a section titled "My Compounds" with three buttons: "Show Structures", "Download SDF", and "Download Smiles". The main content area displays the message "No compounds uploaded." A sidebar on the left contains sections for "WORKBENCH" (with links for "My Compounds" and "Add Compounds") and "TOOLS" (with links for "Past Jobs", "Upload Numeric Data", and "Data").



ChemMine Tools

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ChemMine Tools

Not secure | chemmine.ucr.edu/accounts/signup/

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ChemMine Tools

- WORKBENCH**
 - [My Compounds](#)
 - [Add Compounds](#)
- TOOLS**
 - [Past Jobs](#)
 - [Upload Numeric Data](#)
 - [Cluster](#)
 - [Physicochemical Properties](#)
 - [Similarity Workbench](#)

Signup

Email:

Create password:

Repeat password:



The screenshot shows a web browser window with the address bar displaying "chemmine.ucr.edu/myCompounds/". The page has a dark header with the title "ChemMine Tools" and navigation links for "About", "Help", and "Downloads". Below the header, there is a section titled "My Compounds" with three buttons: "Show Structures", "Download SDF", and "Download Smiles". The main content area states "No compounds uploaded." A sidebar on the left contains sections for "WORKBENCH" (My Compounds, Add Compounds) and "TOOLS" (Past Jobs, Upload Numeric Data). The browser's address bar also shows "Not secure" and various browser icons.



ChemMine Tools

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The screenshot shows a web browser window with the URL `chemmine.ucr.edu/accounts/signin/`. The page title is "ChemMine Tools". On the left, there is a sidebar menu with the following items:

- WORKBENCH**
 - [My Compounds](#)
 - [Add Compounds](#)
- TOOLS**
 - [Past Jobs](#)
 - [Upload Numeric Data](#)
 - [Cluster](#)
 - [Physicochemical Properties](#)
 - [Similarity Workbench](#)

The main content area is titled "Signin" and contains the following form elements:

- Email:
- Password:
- Remember me for a month
-
- [Forgot your password?](#)



ChemMine Tools

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The screenshot shows a web browser window with the URL `chemmine.ucr.edu/accounts/signin/`. The page title is "ChemMine Tools". On the left, there is a sidebar menu with two main sections: "WORKBENCH" and "TOOLS". Under "WORKBENCH", there are links for "My Compounds" and "Add Compounds". Under "TOOLS", there are links for "Past Jobs", "Upload Numeric Data", "Cluster", "Physicochemical Properties", "Similarity", and "Workbench". The main content area is titled "Signin" and contains a form with the following fields and options:

- Email:
- Password:
- Remember me for a month
-
- [Forgot your password?](#)

ChemMine Tools | Morphine - Wikipedia

en.wikipedia.org/wiki/Morphine

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Morphine is used primarily to treat both acute and chronic severe pain. It is also used for pain due to myocardial infarction and for labor pains.^[21] Its duration of analgesia is about three to seven hours.^{[6][7]}

However, concerns exist that morphine may increase mortality in the event of non ST elevation myocardial infarction.^[22] Morphine has also traditionally been used in the treatment of acute pulmonary edema.^[21] A 2006 review, though, found little evidence to support this practice.^[23] A 2016 Cochrane review concluded that morphine is effective in relieving cancer pain. Side-effects of nausea and constipation are rarely severe enough to warrant stopping treatment.^[24]

Shortness of breath [edit]

Morphine is beneficial in reducing the symptom of shortness of breath due to both cancer and noncancer causes.^{[25][26]} In the setting of breathlessness at rest or on minimal exertion from conditions such as advanced cancer or end-stage cardiorespiratory diseases, regular, low-dose sustained-release morphine significantly reduces breathlessness safely, with its benefits maintained over time.^{[27][28]}

Opioid use disorder [edit]

Morphine is also available as a slow-release formulation for opiate substitution therapy (OST) in Austria, Bulgaria, and Slovenia, for addicts who cannot tolerate either methadone or buprenorphine.^[29]

ChEBI [ChEBI:17303](#) ✓

ChEMBL [ChEMBL70](#) ✓

CompTox Dashboard (EPA) [DTXSID9023336](#) ⚙

ECHA InfoCard [100.000.291](#) ⚙

Chemical and physical data

Formula C₁₇H₁₉NO₃

Molar mass 285.34 g/mol g·mol⁻¹

3D model (JSmol) [Interactive image](#) ⚙

Solubility in water HCl & sulf.: 60 mg/mL (20 °C)

MILES [\[hide\]](#)

CN1CC[C@]23C4=C5C=CC(O)=C4O[C@]1H2[C@@H](O)C=C[C@H]3C@H1C5

InChI [\[show\]](#)

[\(verify\)](#)

ChemMine Tools x W Methadone - Wikipedia x +

en.wikipedia.org/wiki/Methadone

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treatment and in the reduction or cessation of heroin use as measured by self-report and urine/hair analysis but did not affect criminal activity or risk of death.^[16]

Treatment of opioid-dependent persons with methadone follows one of two routes: maintenance or detoxification.^[17] Methadone maintenance therapy (MMT) usually takes place in outpatient settings. It is usually prescribed as a single daily dose medication for those who wish to abstain from illicit opioid use. Treatment models for MMT differ. It is not uncommon for treatment recipients to be administered methadone in a specialist clinic, where they are observed for around 15–20 minutes post dosing, to reduce risk of diversion of medication.^[18]

The duration of methadone treatment programs range from a few months to several years. Given opioid dependence is characteristically a chronic relapsing/remitting disorder, MMT may be lifelong. The length of time a person remains in treatment depends on a number of factors. While starting doses may be adjusted based on the amount of opioids reportedly used, most clinical guidelines suggest doses start low (e.g. at doses not exceeding 40 mg daily) and are incremented gradually.^{[19][20]}

Methadone **maintenance** has been shown to reduce the transmission of blood borne viruses associated with opioid injection, such as hepatitis B and C, and/or HIV.^[19] The principal goals of methadone maintenance are to relieve opioid cravings, suppress the abstinence syndrome, and block the euphoric effects associated with opioids.

Chronic methadone dosing will eventually lead to neuroadaptation, characterised by a syndrome of tolerance and withdrawal (dependence). However, when used correctly in treatment, maintenance therapy has been found to be medically safe, non-sedating, and can provide a slow recovery from opioid addiction.^[19] Methadone has been widely used for pregnant women addicted to opioids.^[19]

Opioid detoxification [edit]

Methadone is composed of the 10- and 11-epimers of the opioid. For the treatment of opioid addiction, the use of the 10-epimer is preferred.

CAS Number 76-99-3 [↗](#)

PubChem CID 4095 [↗](#)

IUPHAR/BPS 5458 [↗](#)

DrugBank DB00333 [↗](#)

ChemSpider 3953 [↗](#)

UNII UC6VBETV1Z [↗](#)

KEGG D08195 [↗](#)

ChEBI CHEBI:6807 [↗](#) **X**

ChEMBL ChEMBL651 [↗](#) **✓**

CompTox Dashboard (EPA) DTXSID7023273 [↗](#) **↗**

ECHA InfoCard 100.000.907 [↗](#) **↗**

Chemical and physical data

Formula C₂₁H₂₇NO

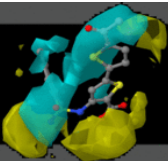
Molar mass 309.445 g/mol g·mol⁻¹

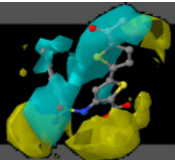
3D model (JSmol) [Interactive image ↗](#)

Chirality Racemic mixture

SMILES [\[hide\]](#)
CC(C)(C1=CC=CC=C1)(C2=CC=CC=C2)CC(N)(C)C

InChI [\[show\]](#)
X (what is this?) [\(verify\)](#)





ChemMine Tools | Methadone - Wikipedia

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ChemMine Tools | About | Help | Downloads | Logged in as rino.ragno@uniroma1.it | Logout

Add Compounds

SMILES Input | Draw a Molecule | SDF Input | SDF Upload | PubChem Import

Enter smiles entry:

```
CN1CC[C@]23C4=C5C=CC(O)=C4O[C@H]2[C@@H](O)C=C[C@H]3[C@H]1C5 Morphine  
CCC(C(C1=CC=CC=C1)(C2=CC=CC=C2)CC(N(C)C)C)=O Methadone
```

Tags

Opioids

Test

Opioids

ChemMine Tools | Methadone - Wikipedia | +

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WORKBENCH

- My Compounds
- Add Compounds

TOOLS

- Past Jobs
- Upload Numeric Data
- Cluster
- Physicochemical Properties
- Similarity Workbench
- Target Search

SEARCH

- Structural Similarity

Success: Added 2 compounds. X

My Compounds

Show Structures | Download SDF | Download Smiles

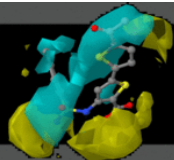
Select all | Deselect all | Add Tags | Remove Tags | Delete Selected

Showing 1 to 2 of 2 entries

Search:

Show 10 entries

CID	Name	Formula	Tags	Options
Methadone		C21H27NO	Test, Opioids, Opioids,	
Morphine		C17H19NO3	Test, Opioids, Opioids,	



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WORKBENCH

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- [Add Compounds](#)

TOOLS

- [Past Jobs](#)
- [Upload Numeric Data](#)
- [Cluster](#)
- [Physicochemical Properties](#)
- [Similarity Workbench](#)
- [Target Search](#)

SEARCH

- [Structural Similarity](#)

Compound Similarity

Select two compounds to compare from the grid below.

Selected Compounds

All

Update

Morphine	Methadone
-----------------	------------------



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ChemMine Tools | Methadone - Wikipedia

chemmine.ucr.edu/similarity/

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remove

All

Update

Morphine

Methadone

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ChemMine Tools x W Methadone - Wikipedia x +

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Apps | Università | WWW | Mail | Ricerca | Vari | SoftwareVari | NatComp | METZ | IMP | Kahoot! | Learning G... | Save to Mendeley

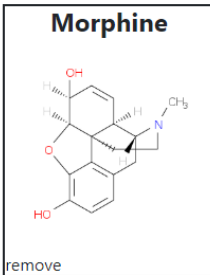
ChemMine Tools About Help Downloads Logged in as rino.ragno@uniroma1.it | Logout

Compound Similarity

Select two compounds to compare from the grid below.

Selected Compounds

Morphine



remove

AP Tanimoto:
0.202597

MCS Tanimoto:
0.5172

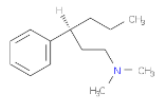
MCS Size: 15

MCS Min: 0.7143

MCS Max: 0.6522

SMILES: [C@@H](c1cccc1)(CCN(C)CCC

Morphine



All x

Update

Morphine

Methadone

WORKBENCH

[My Compounds](#)

[Add Compounds](#)

TOOLS

[Past Jobs](#)

[Upload Numeric Data](#)

[Cluster](#)

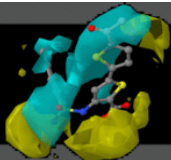
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[Structural Similarity](#)





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