

Fernán Agüero Instituto de Investigaciones Biotecnológicas, UNSAM

# BÚSQUEDA DE SUBESTRUCTURAS/SIMILITUD: FINGERPRINTS

El fingerprint debe ser definido de antemano: bits, folding, count vectors...

Distintas aplicaciones pueden generar distintos tipos de fingerprints

OpenBabel: <a href="https://openbabel.org/docs/dev/Features/Fingerprints.html">https://openbabel.org/docs/dev/Features/Fingerprints.html</a>

```
$ babel -L fingerprints
FP2    Indexes linear fragments up to 7 atoms.
FP3    SMARTS patterns specified in the file patterns.txt
FP4    SMARTS patterns specified in the file SMARTS_InteLigand.txt
MACCS    SMARTS patterns specified in the file MACCS.txt
```

Daylight: https://www.daylight.com/dayhtml/doc/theory/theory.finger.html

RDKit: https://www.rdkit.org/docs/GettingStartedInPython.html#fingerprinting-and-molecular-similarity

# Cuestiones a tener en cuenta

O'Boyle NM, Banck M, James CA, Morley C, Vandermeersch T, Hutchison GR. Open Babel: An open chemical toolbox. J Cheminform. 2011 Oct 7;3:33. doi: 10.1186/1758-2946-3-33. PMID: 21982300; PMCID: PMC3198950.

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# **DETOUR**

Mirar patrones SMARTS en Github

FP3

https://github.com/openbabel/openbabel/blob/master/data/patterns.tx

FP4

https://github.com/openbabel/openbabel/blob/master/data/SMARTS\_InteLigand.txt

**MACCS** 

https://github.com/openbabel/openbabel/blob/master/data/MACCS.txt

# BÚSQUEDA DE SUBESTRUCTURAS

### Screenings

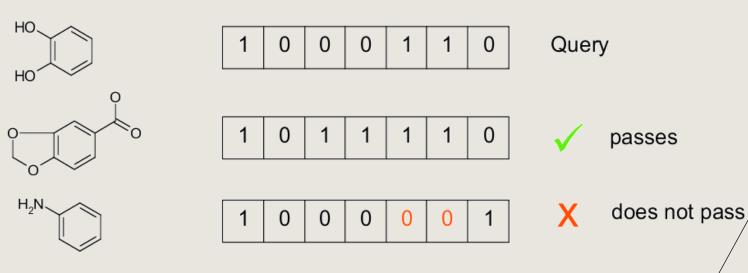
### Simple:

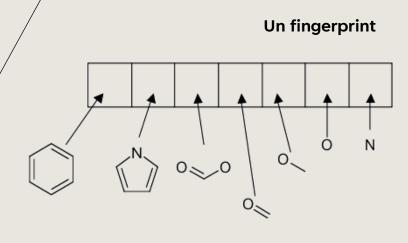
- Usa la fórmula molecular
  - La fórmula de todos los compuestos está almacenada en la base de datos
  - La fórmula de la molécula query se calcula al inicio de la búsqueda
  - Se descartan moléculas a las que les faltan átomos requeridos

# BÚSQUEDA DE SUBESTRUCTURAS: FINGERPRINTS

Fingerprint: representación abstracta de características o propiedades de una molécula (features)

- Presencia/ausencia de cada elemento
- Configuraciones electrónicas inusuales (carbono sp3, nitrógeno unido con un triple enlace)
- Anillos y sistemas de anillos (naftaleno, piridina, cyclohexano)
- Grupos funcionales (alcoholes, aminas, carboxilos, etc.)
- Se suelen utilizar tanto para búsquedas de subestructuras como para detectar similitud



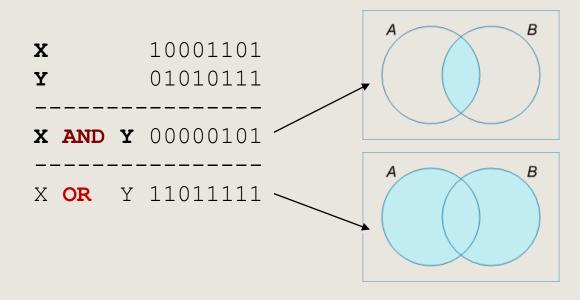


# BÚSQUEDA DE SUBESTRUCTURAS Y SIMILITUD: FINGERPRINTS

Ventajas: screening extremadamente rápido

Se evalúa equivalencia entre conjuntos de bits usando el operador AND binario

Se pueden calcular distancias de similitud a partir de los bits significativos



# DISTANCE METRICS: SIMILARITY, DISIMILARITY

Cociente entre el tamaño de la intersección y el tamaño de la unión de los conjuntos de datos

Jaccard index (J) = Jaccard similarity coefficient = Tanimoto Index = Tanimoto similarity coefficient

(tambien llamado "Intersection Over Union")

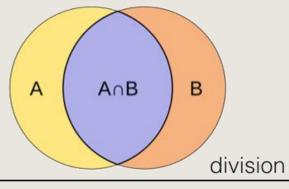
Compara similitudes entre conjuntos de datos finitos

Jaccard distance (d<sub>1</sub>)

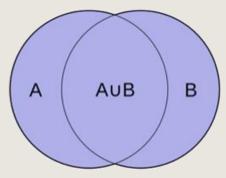
$$J(A,B)=rac{|A\cap B|}{|A\cup B|}=rac{|A\cap B|}{|A|+|B|-|A\cap B|}$$

$$d_J(A,B)=1-J(A,B)=rac{|A\cup B|-|A\cap B|}{|A\cup B|}.$$

The intersect of A & B



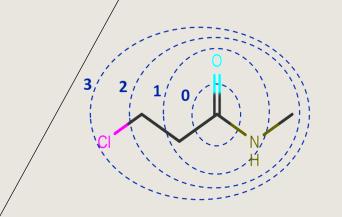
The union of A & B



### Concepto similar al de "extended connectivity" de Morgan

- 1. Assign each atom with an identifier
- 2. Update each atom's identifiers based on its neighbors
- 3. Remove duplicates
- 4. Fold list of identifiers into a 2048-bit vector (a Morgan fingerprint)





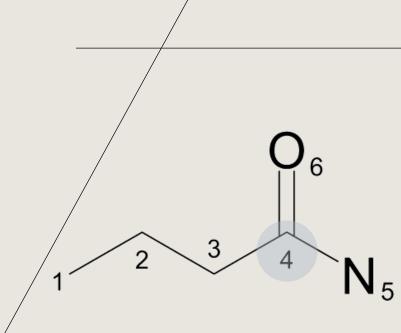
Extended Connectivity
Circular Fingerprints
ECFP6 (radius = 3)
1024 or 2048 bits

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#### 1. Assign each atom with an identifier

We choose an atom in the molecule (e.g. #4) and take note of:

- number of nearest-neighbour non-hydrogen atoms: 3
- number of bonds attached to the atom (not including bonds to hydrogens): 4
- atomic number: 6
- atomic mass: 12
- number of hydrogens connected to the atom: 0
- is the atom in a ring (1) or not (0)?: 0
- Resulting list of numbers is (3,4,6,12,0,0)
- Hash this list of numbers into an integer (identifier)
  - In Python: hash((3, 4, 6, 12, 0, 0, 0))  $\rightarrow$  -5700861834356229464

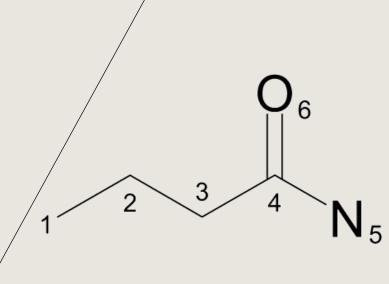


A beginner's guide for understanding Extended-Connectivity Fingerprints (ECFPs). Manish Kumar (2021).

https://chemicbook.com/2021/03/25/a-beginners-guide-forunderstanding-extended-connectivity-fingerprints.html

```
# identificadores para cada atomo
atomo1 = hash((1, 1, 6, 12, 0, 3, 0)) \# -CH3
atomo2 = hash((2, 2, 6, 12, 0, 2, 0)) # -CH2
atomo3 = hash((2, 2, 6, 12, 0, 2, 0)) # -CH2
atomo4 = hash((3, 4, 6, 12, 0, 0, 0)) # -C
atomo5 = hash((1, 2, 7, 14, 0, 0, 0)) \# -NH2
atomo6 = hash((1, 2, 8, 16, 0, 0, 0)) \# = 0
atomo 1 4940186308562569707
atomo 2 -7815985147897826576
atomo 3 -7815985147897826576
atomo 4 -5700861834356229464
atomo 5 -6296387744277800866
```

atomo 6 8618411755682373892



A beginner's guide for understanding Extended-Connectivity Fingerprints (ECFPs). Manish Kumar (2021).

List of

(6)

10

features

https://chemicbook.com/2021/03/25/a-beginners-guide-for-understanding-extended-connectivity-fingerprints.html

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#### Update each atom's identifiers based on its neighbors

Each atom collects its identifier and the identifiers of its immediately neighboring atoms, into an array (list)

And we hash this list again into a new identifier.

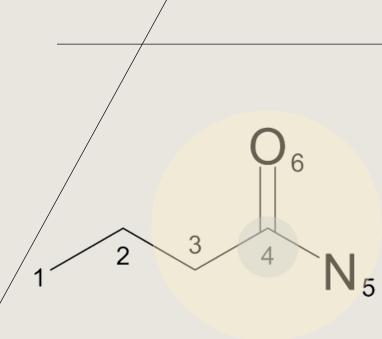
#### Paso anterior

```
atomo 1 4940186308562569707
atomo 2 -7815985147897826576
atomo 3 -7815985147897826576
atomo 4 -5700861834356229464
atomo 5 -6296387744277800866
atomo 6 8618411755682373892
```

```
atomo4_updated = hash((
    1, -5700861834356229464,
    1, -7815985147897826576,
    1, -6296387744277800866,
    2, 8618411755682373892
))

-6784272694619664722

repetimos para los 6 átomos
```



A beginner's guide for understanding Extended-Connectivity Fingerprints (ECFPs). Manish Kumar (2021).

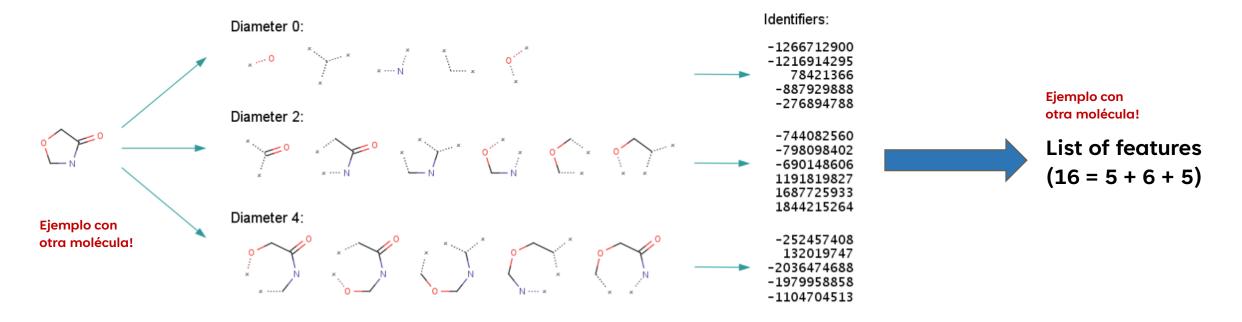
https://chemicbook.com/2021/03/25/a-beginners-guide-for-understanding-extended-connectivity-fingerprints.html

List of

(12)

features

- After that, several iterations are performed to combine the initial atom identifiers with identifiers of neighboring atoms *until a specified diameter is reached*. Each iteration captures larger and larger circular neighborhoods around each atom
- ECFP4 = Extended Circular Fingerprint with radius = 4
- ECFP6 = Extended Circular Fingerprint with radius = 6



## FINGERPRINTS: FOLDING AND BIT COLLISIONS

Para acomodar estos features en un fingerprint de 1024 bits

- Inicializar el fingerprint con todos los bits en 0 (OFF)
- Dividir cada identificador por 1024, y anotar el resto de la división
  - En Python: operador módulo (%)
- Ese es el número de bit 

  que se pone en 1 (ON)

### Resto

### **Ejemplos:**

132019747 % 1024 = 547 1687725933 % 1024 = 877 -798098402 % 1024 = 30**Folding** 

#### **Bit Collision:**

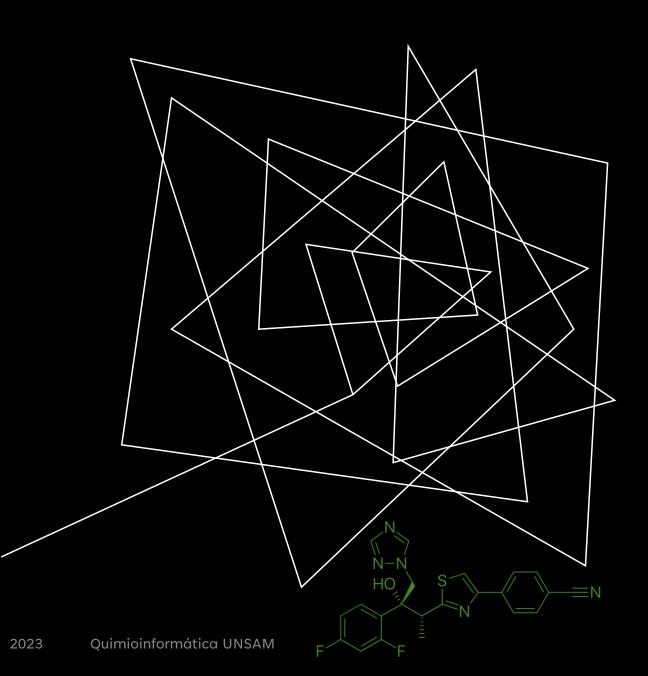
- -14439656419269748 % 1024 = **908**
- -4080868480043360372 % 1024 = **908**

### Solution: increase fingerprint size

- -14439656419269748 % 2048 = 908
- -4080868480043360372 % 2048 = **1932**

**Fixed-length binary representation** 

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# EJEMPLOS

fingerprinting-with-rdkit.ipynb

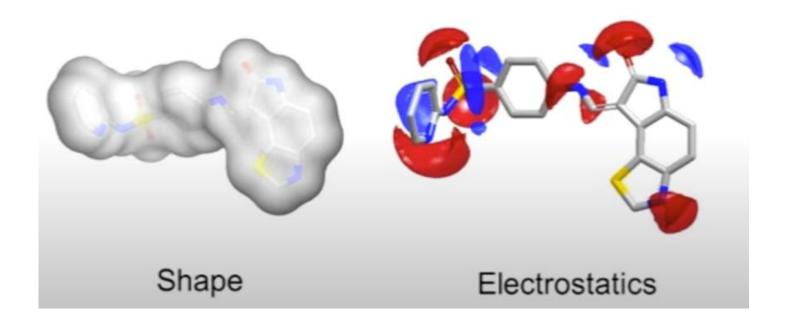
## SCAFFOLD HOPING

computer-aided search for active compounds containing different core structures

Hu Y, Stumpfe D, Bajorath J. Recent Advances in Scaffold Hopping. J Med Chem. 2017 Feb 23;60(4):1238-1246. doi: 10.1021/acs.jmedchem.6b01437. Epub 2016 Dec 21. PMID: 28001064.

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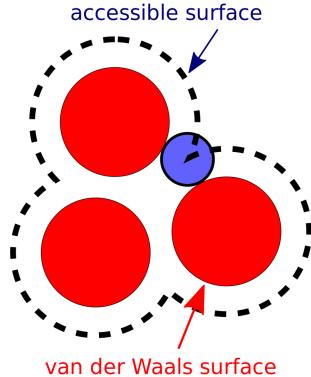
# OTRAS REPRESENTACIONES DE MOLÉCULAS



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## SOLVENT ACCESSIBLE SURFACE AREA CALCULATION

- VSA = van der Waals Surface Area
- AS = Accessible Surface Area



Mitternacht S. FreeSASA: An open source C library for

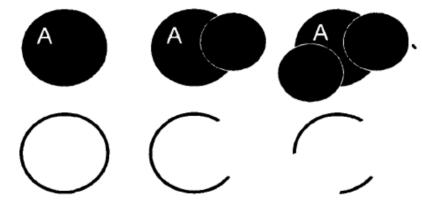


Figure 1. Assuming spherical atoms, the surface area of atom A is the amount of surface area not contained in other atoms.

solvent accessible surface area calculations. F1000Res. 2016 Feb 18;5:189. doi: 10.12688/f1000research.7931.1. PMID: 26973785; PMCID: PMC4776673.

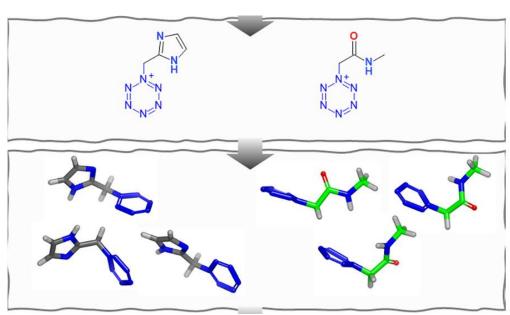
# ESP-SIM: COMPARISON OF ELECTROSTATIC POTENTIAL AND SHAPE

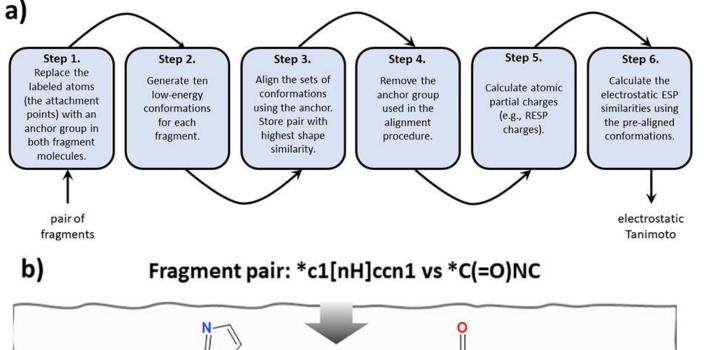
#### https://github.com/hesther/espsim

Bolcato G, Heid E, Boström J. On the Value of Using 3D Shape and Electrostatic Similarities in Deep Generative Methods. J Chem Inf Model. 2022 Mar 28;62(6):1388-1398. doi: 10.1021/acs.jcim.1c01535. Epub 2022 Mar 10. PMID: 35271260; PMCID: PMC8965872.

Step 1. Replace the Instant of PMID: 35271260; PMCID: PMC8965872.

Step 2. Step 3. Align the sets of Remove the Instant of PMID: 35271260; PMCID: PMC8965872.

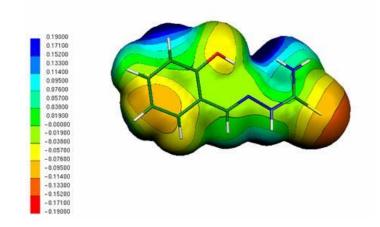


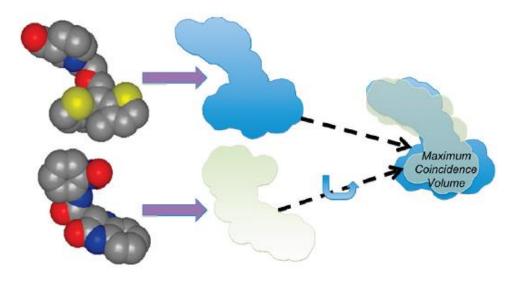


https://github.com/hesther/espsim/blob/master/scripts/short\_demonstration.ipynb

# REPRESENTACIÓN DE MOLECULAS: 3D

- Una representación tridimensional de la molécula requiere no sólo especificar coordenadas espaciales de átomos
  - También hay que especificar
    - Volumen
      - Fused spheres
      - Atom-centered Gaussians
    - Superficie
    - Forma
      - Coincidencia de volumenes



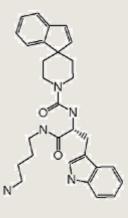


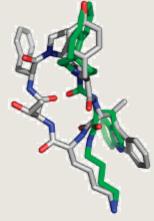
Molecular shape and medicinal chemistry: a perspective. 2010. A Nicholls *et al.* J Med Chem 53: 3862

# REPRESENTACIÓN DE FORMA (SHAPE)

Varias aplicaciones posibles:

- •Búsqueda de moléculas similares
  - •En este caso la similitud es a nivel de forma
    - •Se pueden agregar adicionalmente limitaciones
  - •Varias implementaciones en la industria farmacéutica
    - Virtual screening
      - Varios casos de éxito conocidos
      - •Merck, primer aplicación publicada del método
        - •Identificación de análogos no-peptídicos de:
          - antagonista endógeno del receptor de fibrinógeno (Arg-Gly-Pro)
          - Somatotrophin release inhibitor factor





# REPRESENTACIÓN DE FORMA (SHAPE)

Varias aplicaciones posibles:

### **Lead optimization**

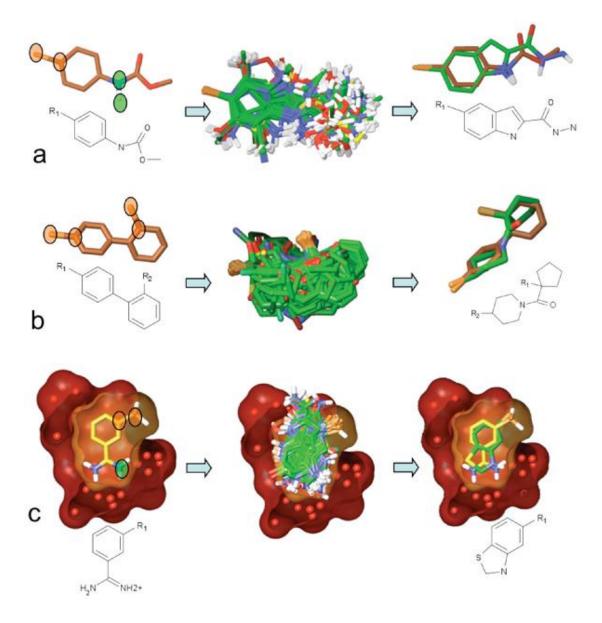
Uno cuenta con una molécula activa que quiere optimizar

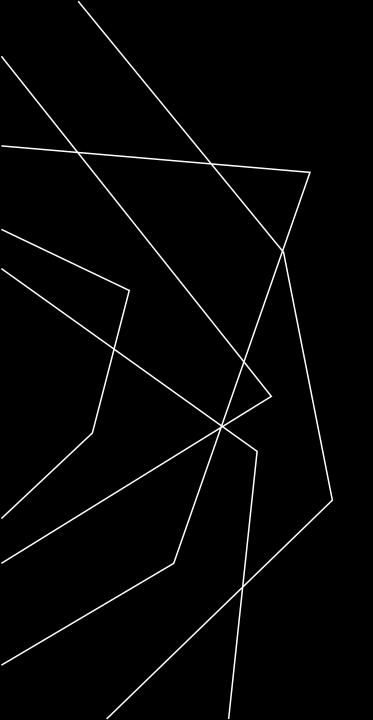
### **Scaffold Hoping**

Facilmente explorable utilizando metodos computacionales

Molecular shape and medicinal chemistry: a perspective. 2010. A Nicholls *et al.* J Med Chem 53: 3862

KIN: Bristol-Myers Squibb





# **INTERVALO**

15 minutos



## CALCULO DE PROPIEDADES

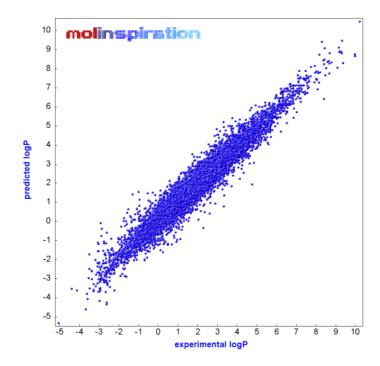
**Enlaces rotables** 

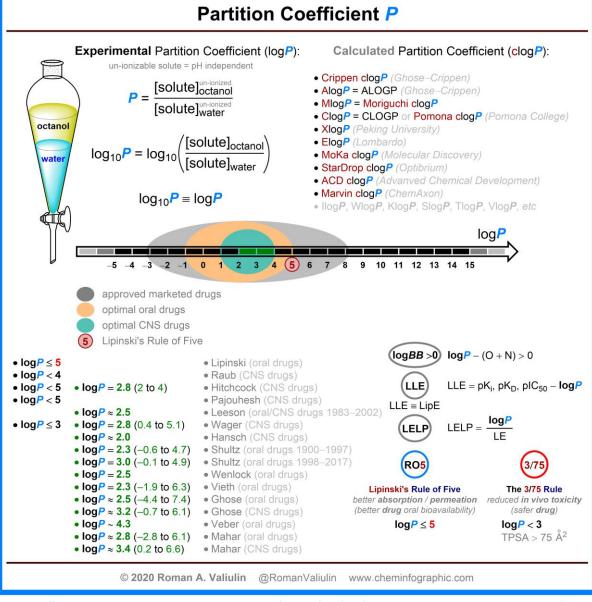
Dadores / Aceptores de puentes de hidrógeno

cLogP (coeficiente de partición octanol / agua)

PSA (polar surface area) / TPSA (topological surface area)

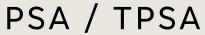
### LOGP PARTITION COEFFICIENT

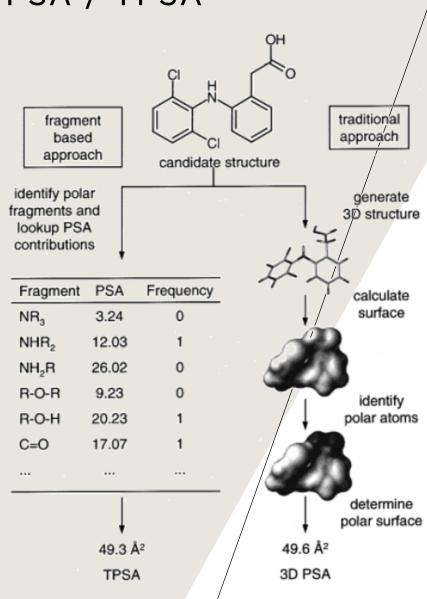


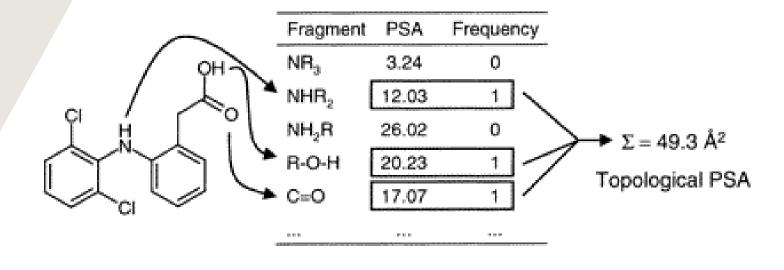


https://cheminfographic.wordpress.com/2020/05/01/partition-coefficient-p-logp/

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### Polar Surface Area (PSA, costoso)

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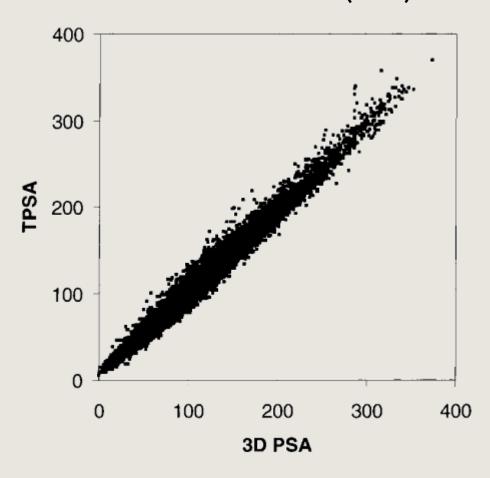
Requiere generar conformeros 3D para calcular SA (Surface Area)

### Topological Polar Surface Area (TPSA)

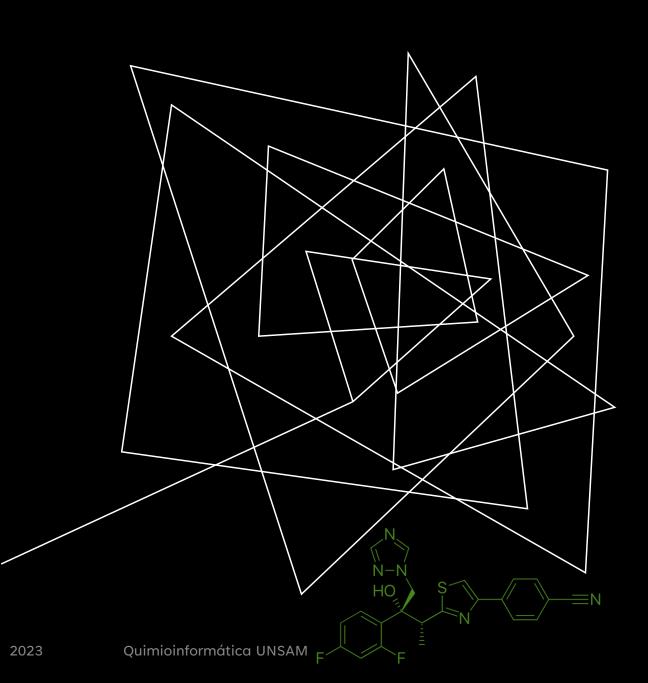
Sumatoria de contribuciones tabuladas de fragmentos polares

Ertl P, Rohde B, Selzer P. Fast calculation of molecular polar surface area as a sum of fragment-based contributions and its application to the prediction of drug transport properties. J Med Chem. 2000 Oct 5;43(20):3714-7. doi: 10.1021/jm000942e. PMID: 11020286.

# TPSA VS PSA (3D)



Ertl P, Rohde B, Selzer P. Fast calculation of molecular polar surface area as a sum of fragment-based contributions and its application to the prediction of drug transport properties. J Med Chem. 2000 Oct 5;43(20):3714-7. doi: 10.1021/jm000942e. PMID: 11020286.



# ONTOLOGIAS

Organizando datos químicos

# QUE ES UNA ONTOLOGIA?

Es una formalización de un área del conocimiento mediante reglas

# QUÉ SIGNIFICA ONTOLOGÍA?

### **Webster's Revised Unabridged Dictionary**

### Ontology

- the things which exist
- The department of the science of metaphysics which investigates and explains the nature and essential properties and relations of all beings,

### The Free On-Line Dictionary of Computing

### Ontology

- Phylosophy: a systematic account of experience
- Artificial Inteligence: an explicit formal specification of how to represent the objects, concepts and other entities that are assumed to exist in some area of interest and the relationships that hold among them. [...]
- Information Science: the hierarchical structuring of knowledge about things by subcategorizing them according to their essential (or at least relevant and/or cognitive) qualities.

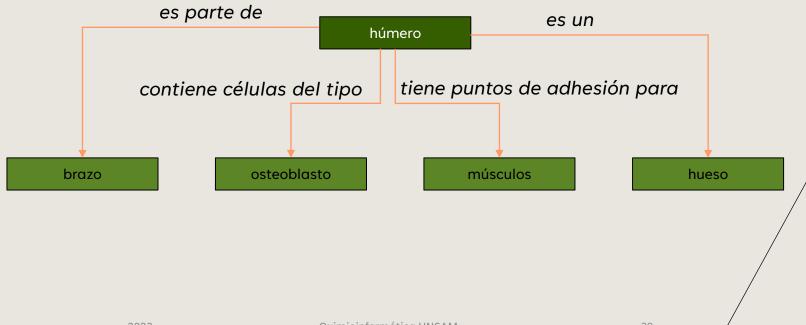
### EJEMPLO: UNA ONTOLOGIA ANATOMICA

### Una ontología es un área del conocimiento que ha sido formalizada:

- **Términos** (conceptos) individuales
- Afirmaciones (reglas) que conectan términos entre sí

### Ejemplo: una ontología anatómica:

- Términos: húmero, brazo, osteoblasto, músculo, hueso
- Conexiones: es parte de, contiene células del tipo, tiene puntos de adhesión para, es un



2023

## ONTOLOGIAS QUIMICAS

ChEBI – Chemical Entities of Biological Interest <a href="https://www.ebi.ac.uk/chebi/">https://www.ebi.ac.uk/chebi/</a>

Son varias ontologías agrupadas:

- Molecular Structure
  - Términos: hydrocarbons, carboxylic acids, tertiary amines
  - Afirmaciones: (reglas): is a (es un), has part (contiene)

caffeine (CHEBI:27732) **is a** purine alkaloid (CHEBI:26385) caffeine (CHEBI:27732) **is a** trimethylxanthine (CHEBI:27134)

caffeine monohydrate (CHEBI:31332) has part caffeine (CHEBI:27732) sodium caffeine benzoate (CHEBI:32140) has part caffeine (CHEBI:27732)

## ONTOLOGIAS QUIMICAS

ChEBI – Chemical Entities of Biological Interest

https://www.ebi.ac.uk/chebi/

Son varias ontologías agrupadas:

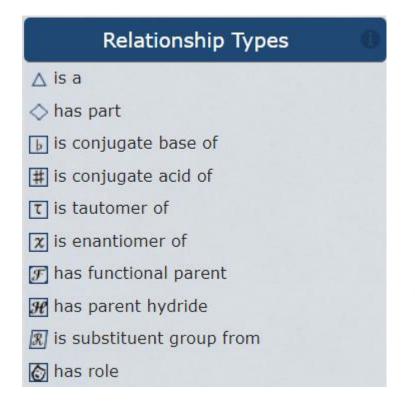
- Role
  - Chemical Role
    - ej ligand, inhibitor, surfactant
  - Biological Role
    - antibiotic, antiviral agent, coenzyme, hormone
  - Application
    - pesticide, antirheumatic drug, fuel
  - **Términos:** por ej alguna entidad molecular (o una parte)
  - Afirmaciones: (reglas): has role (tiene rol de)

caffeine (CHEBI:27732) has role environmental contaminant (CHEBI:78298)

caffeine (CHEBI:27732) has role adenosine A2A receptor antagonist (CHEBI:53121)

caffeine (CHEBI:27732) has role food additive (CHEBI:64047)

# CHEBI: AFIRMACIONES (REGLAS)



△ CHEBI:17026 progesterone

TCHEBI:16973 11-deoxycorticosterone

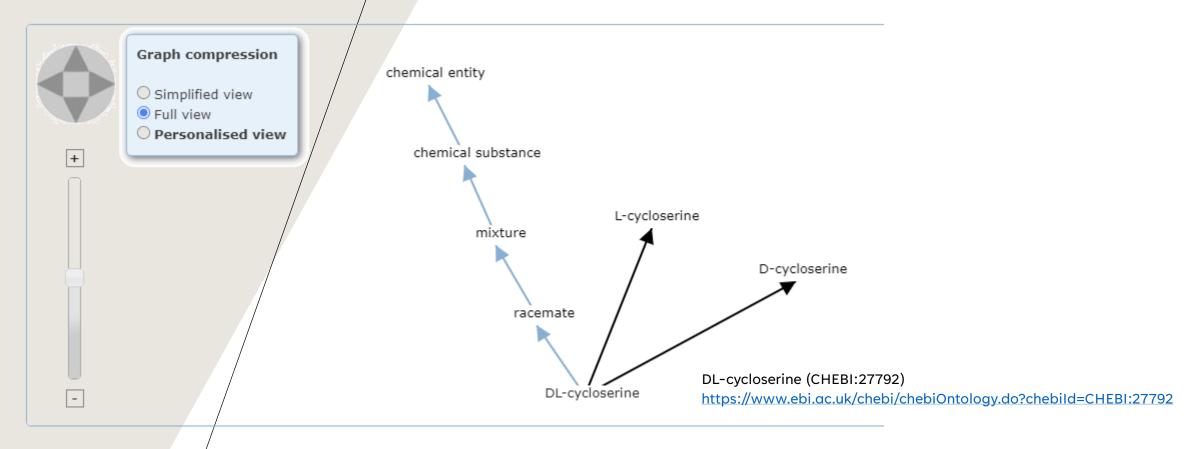
# CHEBI: AFIRMACIONES (REGLAS)

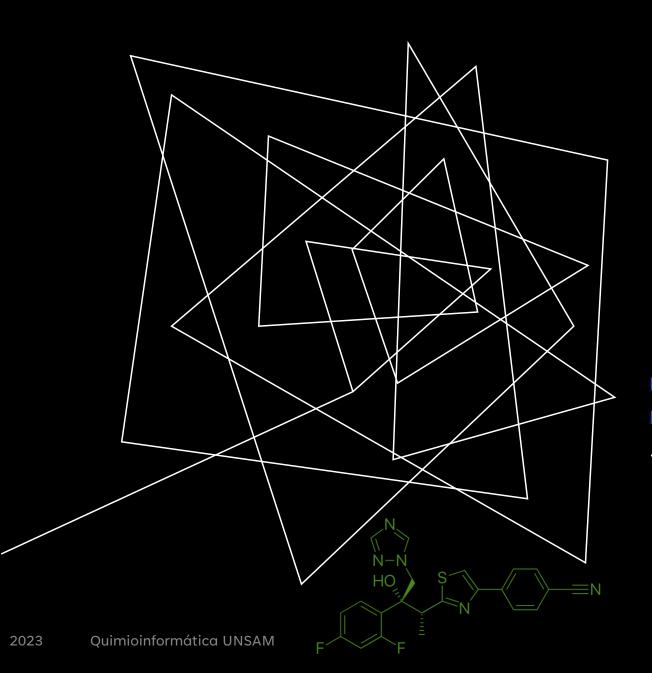
### Relationship Types

- △ is a
- has part
- ы is conjugate base of
- # is conjugate acid of
- T is tautomer of
- x is enantiomer of
- F has functional parent
- # has parent hydride
- IN is substituent group from
- has role

- △ CHEBI:16482 naphthalene
- CHEBI:50715 methylnaphthalene
- △ CHEBI:16414 L-valine
- CHEBI:32853 L-valyl group
- △ CHEBI:35482 opioid analgesic
- CHEBI:17303 morphine

# NAVEGACIÓN GRAFICA DE LA ONTOLOGIA

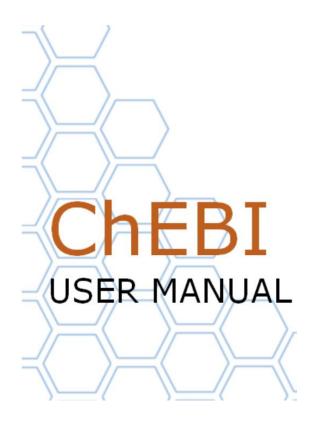




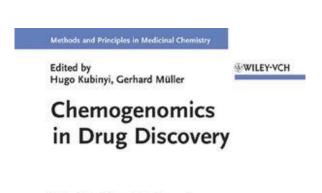
# **EJEMPLOS**

https://www.ebi.ac.uk/chebi/searchId.do?chebiId=CHEBI:27732 https://www.ebi.ac.uk/chebi/searchId.do?chebiId=CHEBI:78298

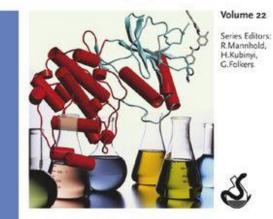
# BIBLIOGRAFÍA | MATERIAL DE LECTURA



https://www.ebi.ac.uk/chebi/aboutChebi Forward.do

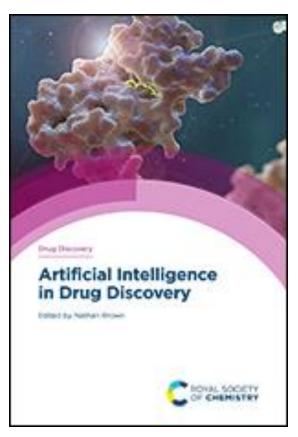


A Medicinal Chemistry Perspective



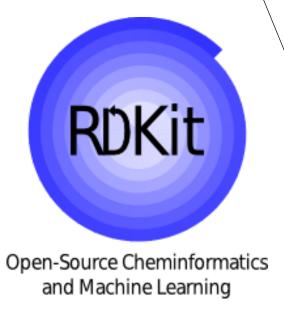
Chemogenomics in Drug Discovery: A Medicinal Chemistry Perspective (2006). Edited by Hugo Kubinyi & Gerhard Müller, Wiley-VCH.

https://www.wiley.com/enus/Chemogenomics+in+Drug+Discovery%3A +A+Medicinal+Chemistry+Perspective-p-9783527604029



Artificial Intelligence in Drug Discovery (2020). Edited by Nathan Brown. Royal Society of Chemistry.

https://doi.org/10.1039/9781788016841



The RDKit Book (2023).
https://www.rdkit.org/docs/RDKit\_Book.html