

Cheminformatics Introduction

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Associate Instructor

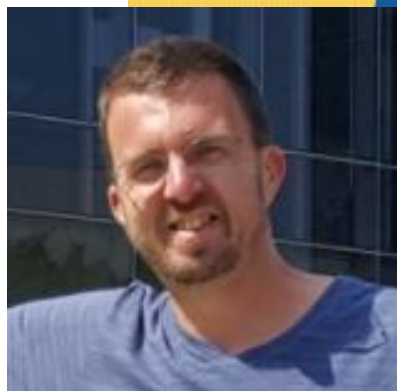
IU School of Informatics and Computing

Instructor: Prof. Joanne Luciano



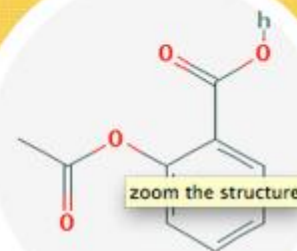
DATA SCIENCE FOR DRUG DISCOVERY, HEALTH AND
TRANSLATIONAL MEDICINE (DSDHT) INFO I-590

Cheminformatics: Introduction



FREE ONLINE COURSE
introducing cheminformatics

learncheminformatics.com



learn cheminformatics!

Cheminformatics is the study of all aspects of the representation and use of chemical and related biological information on computer. It has applications in drug discovery, health, data mining and many other areas

This site gives you links to some resources to get you started in learning about cheminformatics - a free online course, a low-cost eBook, an established open access journal and a

eBook just \$9.95

*Kindle and PDF format
library versions available*

Introducing
cheminformatics

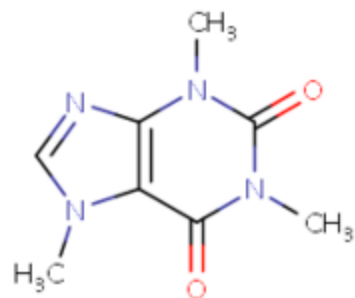
**Introducing
Cheminformatics**
An intensive self-study guide

David J. Wild

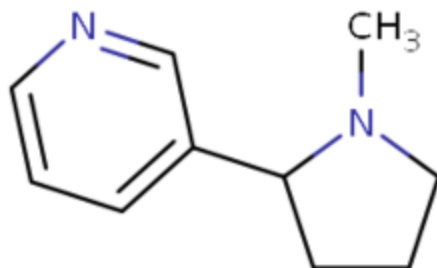
Cheminformatics is the study of all aspects of the representation and use of chemical and related biological information on computer. It has applications in drug discovery, health, data mining and many other areas

Cheminformatics:

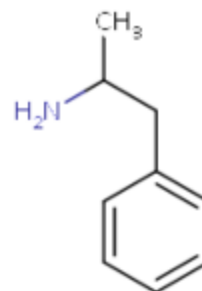
What makes drugs so special?



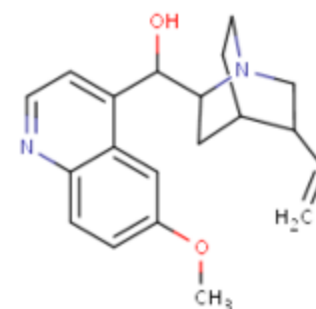
caffeine



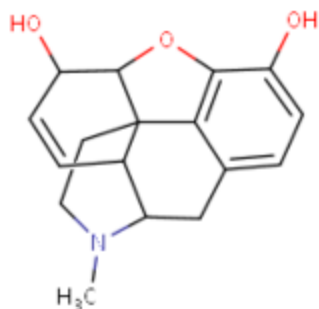
nicotine



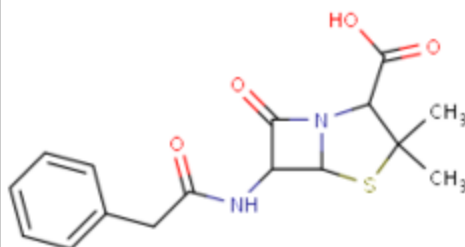
adderall



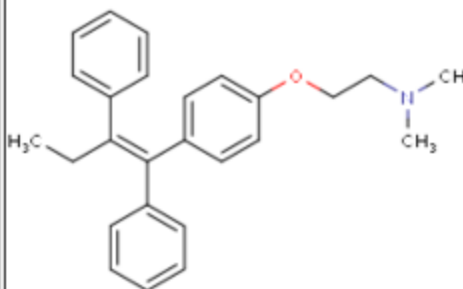
quinine



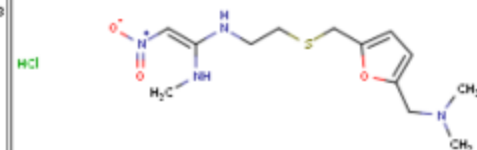
morphine



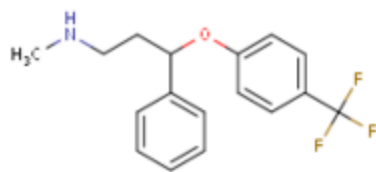
benzylpenicillin



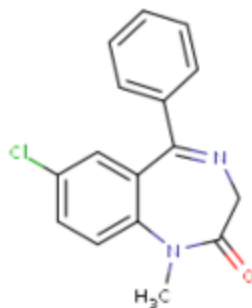
Tamoxifen



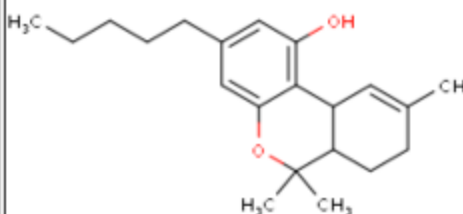
Zantac



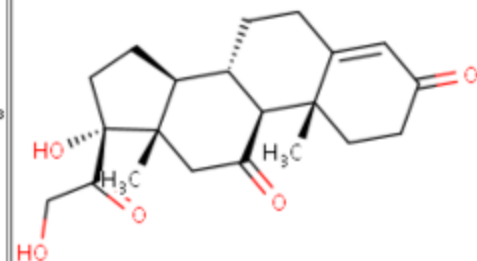
Prozac



Valium



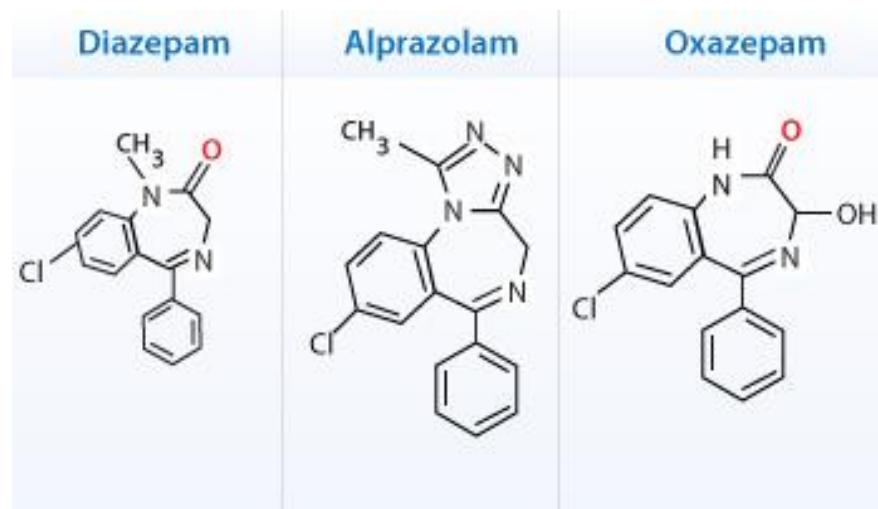
THC



Cortisone

Cheminformatics: Chemotypes

- Benzodiazepines class of tranquilizers sharing a common ring system, a pattern, chemotype or scaffold.
- Similarity principle: similar chemicals, bioactivity.
- Representing chemicals as molecular graphs enables efficient storage and analysis.



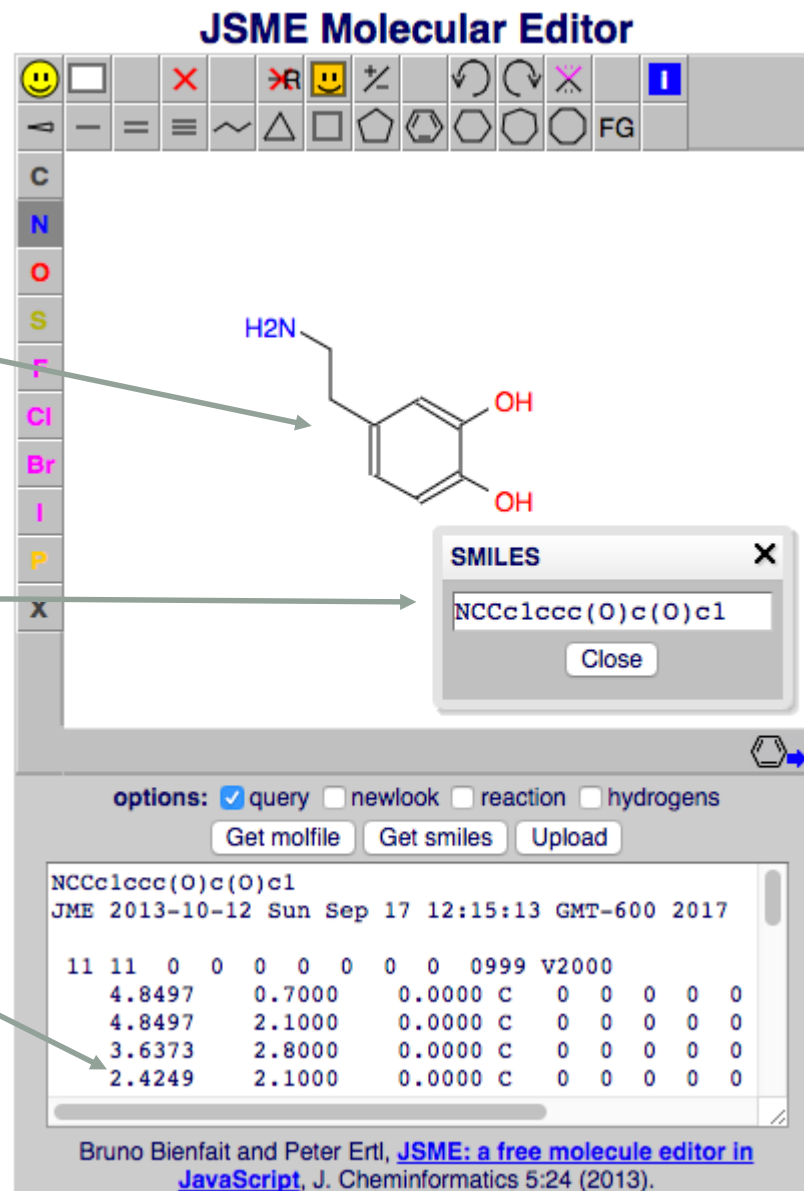
Drug name	Brand names
diazepam	Valium, Ducene, Antenex, Valpam
oxazepam	Serepax, Murelax, Alepam
nitrazepam	Mogadon, Alodorm
temazepam	Normison, Temaze, Temtabs
lorazepam	Ativan
flunitrazepam	Rohypnol, Hypnodorm
bromazepam	Lexotan
clonazepam	Rivotril, Paxam

Cheminformatics: Representing molecules computationally

molecular depiction (2D)

SMILES (Simplified Molecular
Input Line Entry System)

molfile (connection table)



Cheminformatics: Chemical databases

NIH U.S. National Library of Medicine National Center for Biotechnology Information

PubChem | OPEN
CHEMISTRY
DATABASE

Search Compounds

Compound Summary for CID 3016

Download

Share



Diazepam



STRUCTURE



VENDORS



DRUG INFO



PHARMACOLOGY



LITERATURE



PATENTS



BIOACTIVITIES

PubChem CID:

3016

Chemical Names:

Diazepam; Valium; Diazemuls; Relanium; Ansiolesina; Stesolid

More

Molecular Formula:

$C_{16}H_{13}ClN_2O$

Molecular Weight:

284.743 g/mol

InChI Key:

AAOVKJBEBIDNHE-UHFFFAOYSA-N

Drug Information:

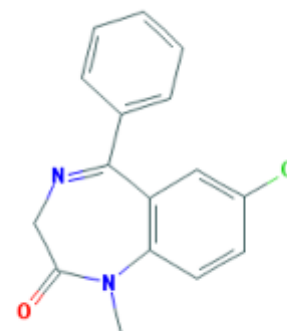
Drug Indication

Therapeutic Uses

Clinical Trials

FDA Orange Book

FDA UNII



"PubChem, released in 2004, provides information on the biological activities of small molecules."

<https://pubchem.ncbi.nlm.nih.gov/>

Cheminformatics: Chemical patterns

Depict

format: SMILES ☒ file2txt

upload: Choose File No file chosen ...or paste:

```
[nH]1c2ccccc2nc1C(c3ccccc3)O NSC 405
n1c(c2[nH]cnc2nc1N)N NSC 743
n1c(c2[nH]cnc2nc1)Cl NSC 744
n1c(=S)c2[nH]cnc2[nH]c1N NSC 752
n1cc2[nH]cnc2nc1 NSC 753
n1c(=S)c2[nH]cnc2[nH]c1 NSC 755
c1c2c(ccc1)nc[nH]2 NSC 759
c1(-O)c2c(n(c(-O)n1C)C)nc[nH]2 NSC 2066
```

options:

- ☐ show Hs
- ☐ lonepairs
- ☒ use2d
- ☒ transparent
- ☒ zoomable
- ☐ verbose
- ☐ show properties
- ☐ show atom maps
- ☐ smilesmatch
- ☐ parts2mols
- ☐ showarom
- ☐ align2smarts

smarts: [#6]12~[#7]~*~[#7]~[#6]1~*~*~*~*2

output:

size: m - 180x260 ncols: 4 mode: COW

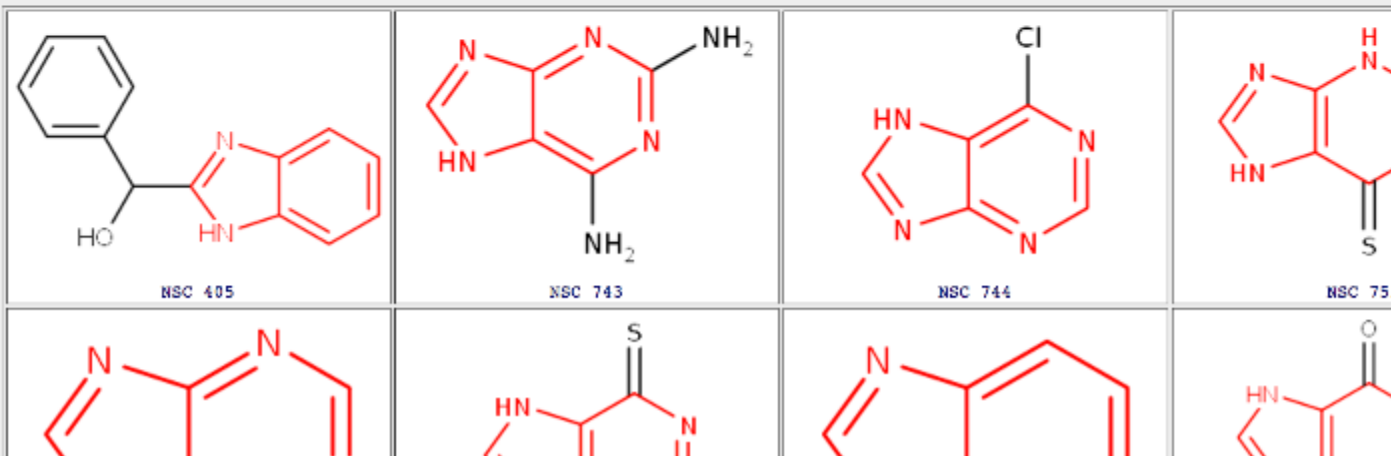
format: ☒ png ☐ jpeg

Go Depict

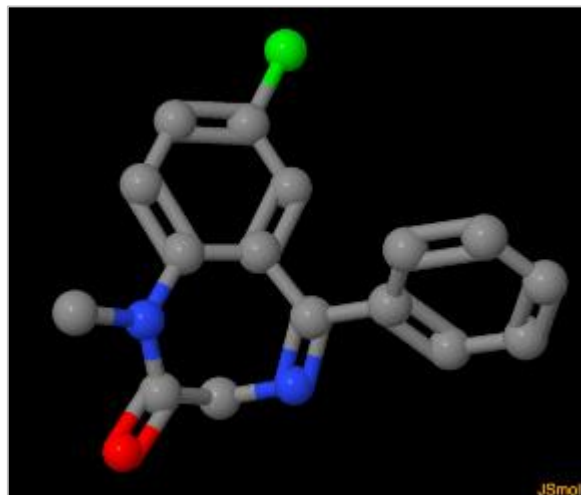
[#6]12~[#7]~*~[#7]~[#6]1~*~*~*~*2

Imidazole pattern
subgraph
matching with
SMARTS.

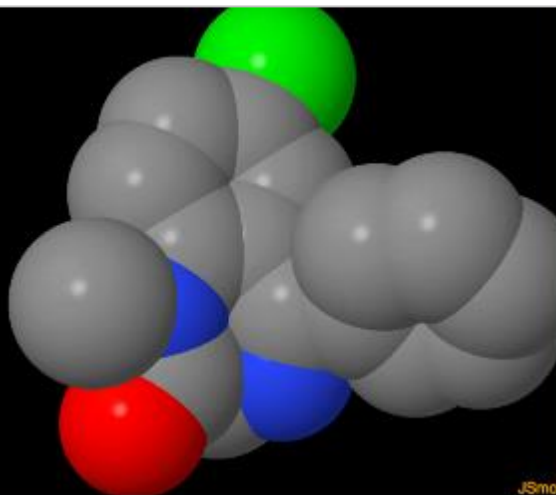
SMARTS =
Smiles Arbitrary
Target
Specification
(akin to regexp)



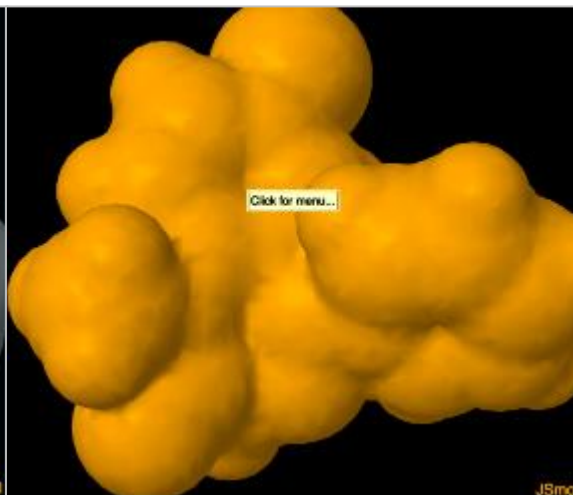
Cheminformatics: Molecules in 3D



ball and stick



spacefill



VDW surface

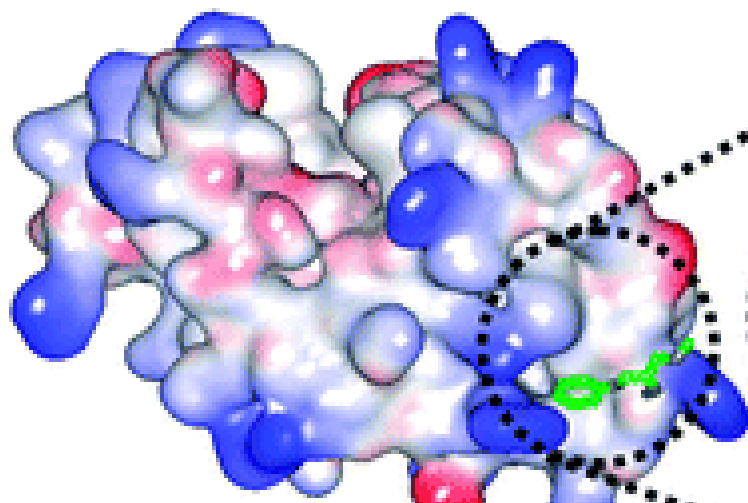
- Drug bioactivity is related to 3D properties and interactions.

Cheminformatics: Glossary

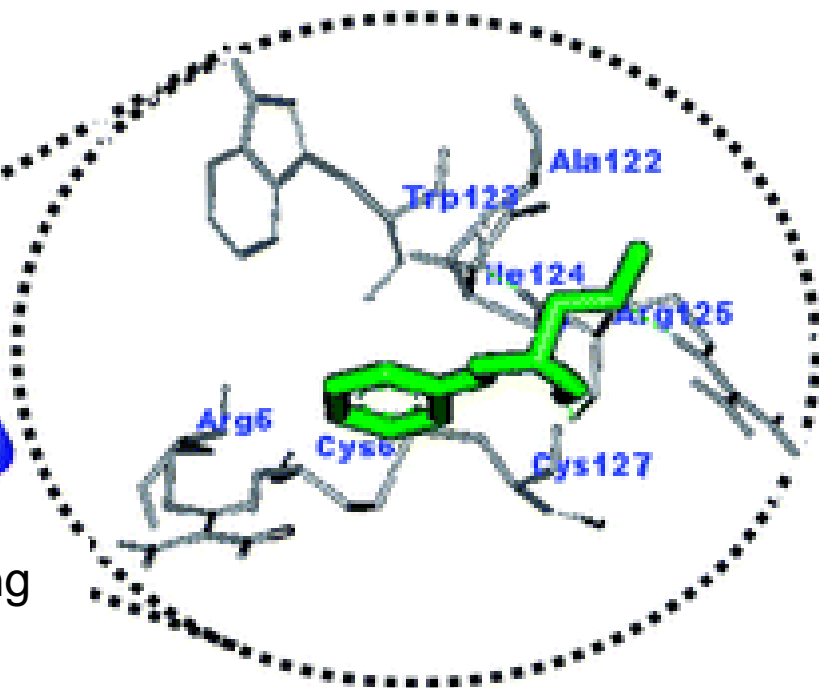
small molecule	not a protein or nucleic acid (macromolecule)
pharmacology	study of uses, effects and modes of action of drugs
bioactivity	effect of a chemical on a biological target
IC50, EC50	measures of bioactive potency
structure-activity relationship	bioactivity vs. structural features within a chemical series
molecular descriptor	measured, theoretical or computed property
substructure searching	searching for molecular subgraphs
similarity searching	searching for similar molecules
pharmacophore	3D features pattern interacting with a binding site
molecular modeling	computational models, 2D, 3D and statistical, for explaining, visualizing and predicting
computational chemistry	connotes physics-based modeling, molecular mechanics and dynamics, etc.
virtual screening	<i>in silico</i> search (assay) for bioactive candidates

Cheminformatics:

Modeling protein-ligand binding



AutoDock is a suite of automated docking tools. It is designed to predict how small molecules, such as substrates or drug candidates, bind to a receptor of known 3D structure.



AutoDock is one of several tools for docking and modeling protein-ligand binding.

Cheminformatics:

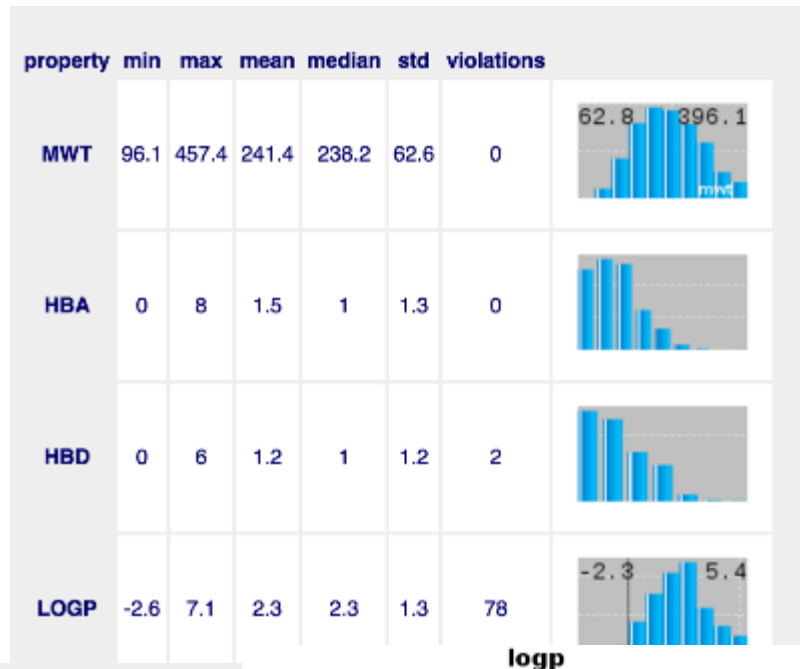
Drug-likeness and lead-likeness

The Lipinski Rule of Five (Ro5) was formulated by Christopher A. Lipinski and published in 1997, and is intended to guide library selection for drug discovery. The following molecular descriptors are used:

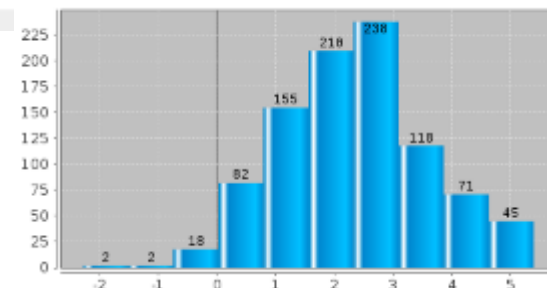
- mwt - avg molecular weight
- hbd - hydrogen bond donor count
- hba - hydrogen bond acceptor count
- LogP - octanol-water partition coefficient

The Ro5 specifies that compounds should violate no more than one of the following criteria:

- mwt ≤ 500
- hbd ≤ 5
- hba ≤ 10
- LogP < 5.0



Ro5 violations	#mols
0	921
1	78
2	1
3	0
4	0



Cheminformatics:

Drug discovery applications

- Databases of relevant compounds and metadata
- 2D virtual screening (substructure, similarity)
- 3D virtual screening (shape, pharmacophores)
- Property prediction (e.g. solubility, logP)
- ADMET prediction (absorption, distribution, metabolism, excretion, toxicity)
- Clustering (unsupervised ML)
- Quantitative structure-activity relationship modeling (QSAR)
- Modeling/predicting protein-ligand binding
- *De novo* design (e.g. GAs)

Cheminformatics:

Take home messages

- Understanding the structure and properties of chemicals enables us to explore biological effects and discover new drugs.
- Chemical databases provide access to a wealth of biomedical and drug discovery data, searchable via cheminformatics.
- In silico predictive modeling accelerates and economizes early stage drug discovery.