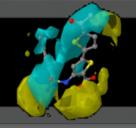


Introduction to Ligand-Based Drug Design



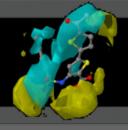


Summary



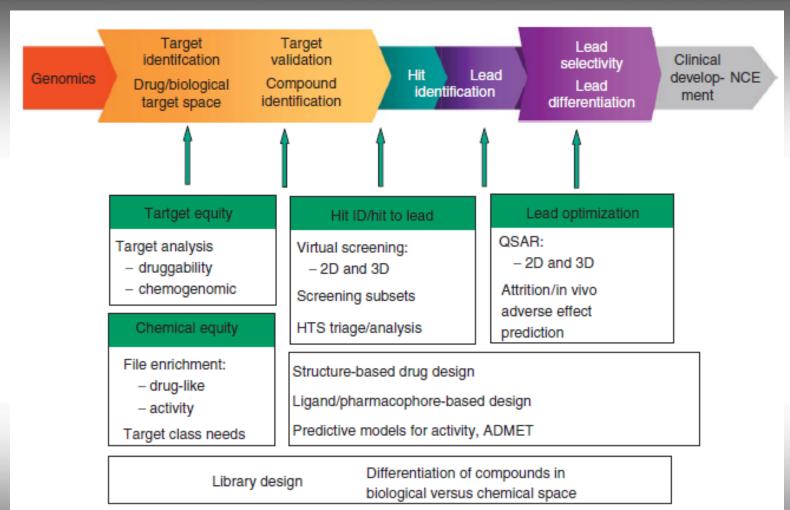
1. Overview on LBDD

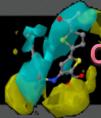
- A. Definition and Methods
- B. Pharmacophore Approach
- C. QSAR
- D.3-D QSAR
- 2. Application (and demonstration) of 3-D QSAR to medicinal chemistry



Drug Discovery Process

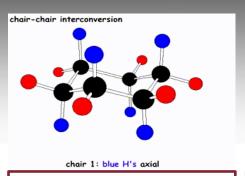


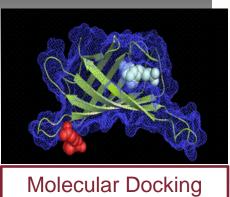


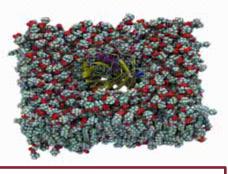


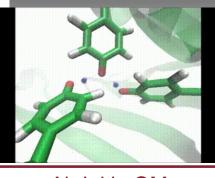
Computational Techniques in Medicinal Chemistry











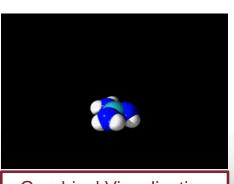
Conformational Search

Simulate Annealing

Ab Initio QM



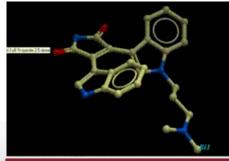
Molecular Dynamics



Graphical Visualization



3-D QSAR



Pharmacophore

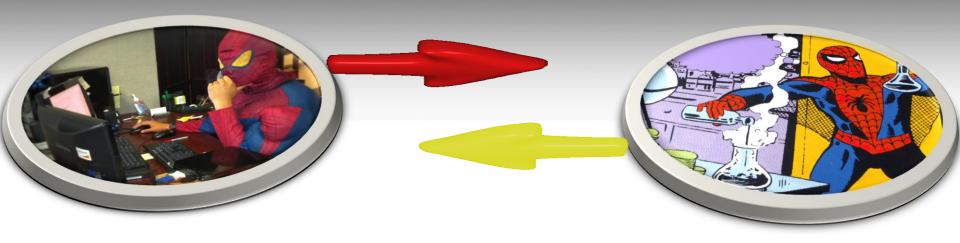
QSAR, COMBINE, Scoring Functions, Homology Modeling,....

QSAR



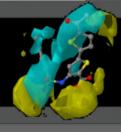


Drug Design = Computational & Synthesis Tandem



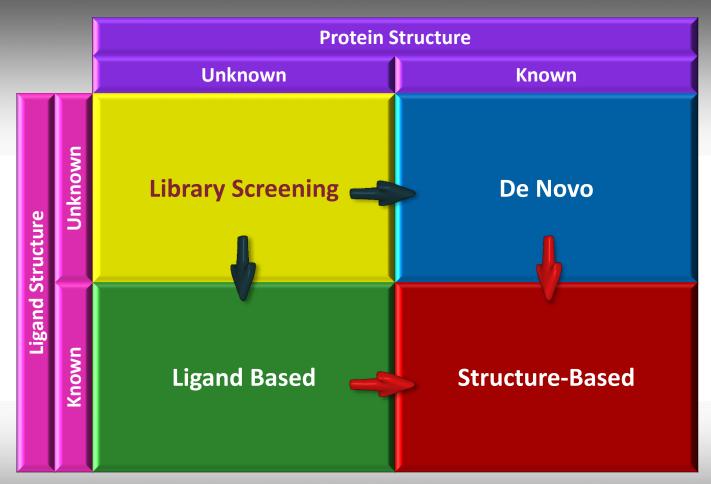


Basic Hit Hit to Lead Clinical **Target Target** Preclin **Prod Diagnos** Research Ident Valid **Ident** Lead Opt **Trial**

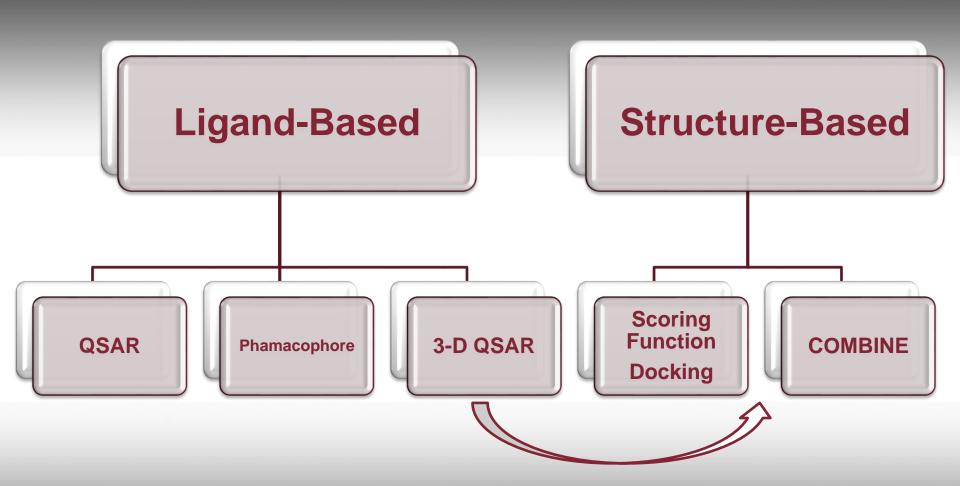


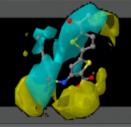
Computational Medicinal Chemistry





6

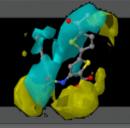




LBDD



In the absence of three-dimensional (3-D) structures of potential drug targets, ligand-based drug design is one of the popular approaches for drug discovery and lead optimization. 3-D structure activity relationships (3-D QSAR) and pharmacophore modeling are the most important and widely used tools in ligandbased drug design that can provide crucial insights into the nature of the interactions between drug target and ligand molecule and provide predictive models suitable for lead compound optimization.

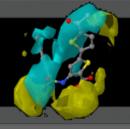


Pharmacophore Definition



"A pharmacophore is the ensemble of steric and electronic features that is necessary to ensure the optimal supramolecular interactions with a specific biological target and to trigger (or block) its biological response."

C.-G. Wermuth et al., *Pure Appl. Chem.* 1998, 70: 1129-1143



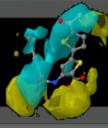
Definition by IUPAC



A **pharmacophore** is the ensemble of steric and electronic features that is necessary to ensure the optimal supramolecular interactions with a specific biological target structure and to trigger (or to block) its biological response.

A **pharmacophore** does not represent a real molecule or a real association of functional groups, but a purely abstract concept that accounts for the common molecular interaction capacities of a group of compounds towards their target structure.

The **pharmacophore** can be considered as the largest common denominator shared by a set of active molecules. This definition discards a misuse often found in the **medicinal chemistry** literature which consists of naming as **pharmacophores** simple chemical functionalities such as guanidines, sulfonamides or dihydroimidazoles (formerly imidazolines), or typical structural skeletons such as flavones, phenothiazines, prostaglandins or steroids.



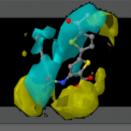
Pharmacophoric Descriptors



Interaction sites used to define a pharmacophore,

- H-bonding,
- hydrophobic
- Electrostatic

Defined by atoms, ring centers and virtual points.



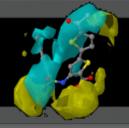
The First Paper on Pharmacophoric Approach



The Conformational Parameter in Drug Design: The Active Analog Approach

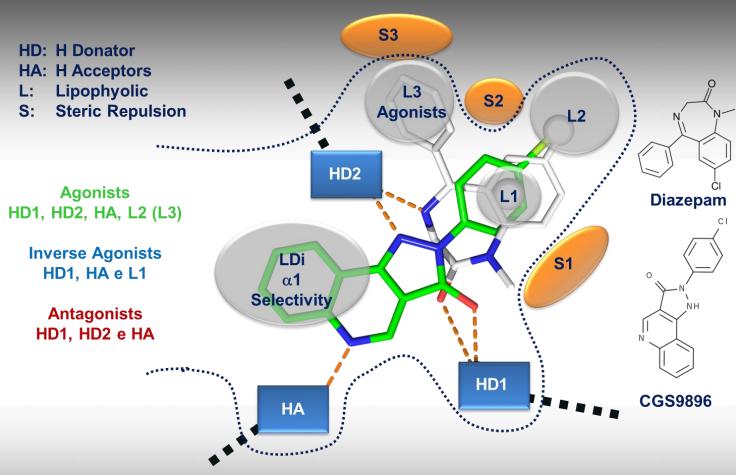
GARLAND R. MARSHALL, C. DAVID BARRY, HEINZ E. BOSSHARD, BICHARD A. DAMMKOEHLER, and DEBORAH A. DUNN

Departments of Physiology and Biophysics, of Pharmacology and of Computer Science and the Computer Systems Laboratory, Washington University, St. Louis, MO 63110

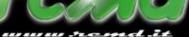


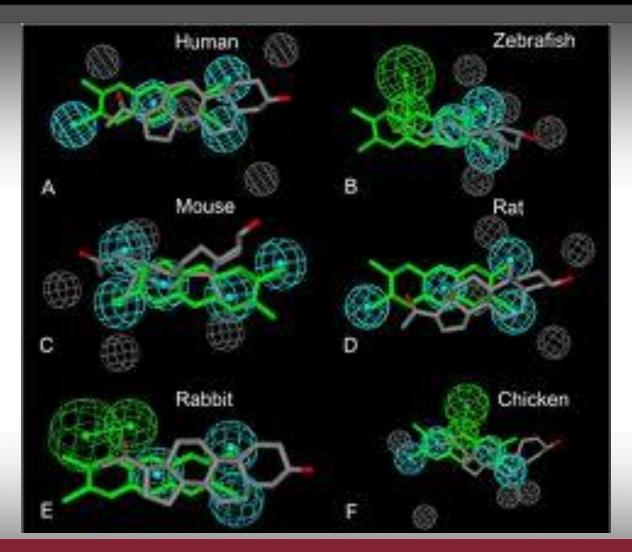
BDZ Cook Model

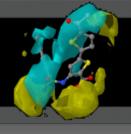




Modern Pharmacophore Model







The Beginning of QSAR

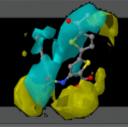


QSAR date back to the 19th century

In 1863, A.F.A. Cros at the University of Strasbourg observed that toxicity of alcohols to mammals increased as the water solubility of the alcohols decreased

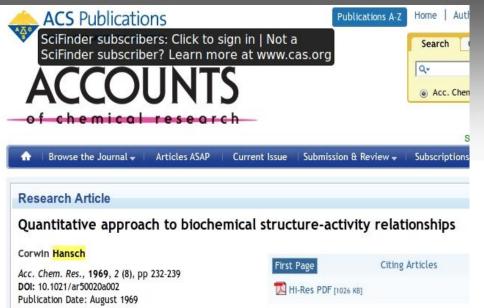
In the 1890's, Hans Horst Meyer of the University of Marburg and Charles Ernest Overton of the University of Zurich, working independently, noted that the toxicity of organic compounds depended on their lipophilicity.

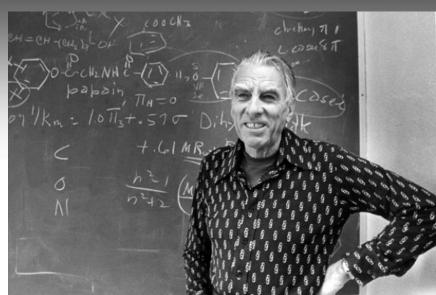
Louis Hammett (1938) Electronic Parameters (Sigma-Rho) Robert W. Taft (1952) Steric Parameters (Es) Corvin Hansch (1964) Hydrophobic Parameter (LogP - π)



The Hansch Equation



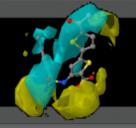




The Hansch Equation

$$\delta_{\rm X} \log k_i = \delta_{\rm X} \Delta G_{\rm hydrophobic} +$$

$$\delta_{\rm X} \Delta G_{\rm electronic} + \delta_{\rm X} \Delta G_{\rm steric}$$





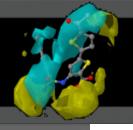
The first QSAR equations were based on the observation that partition coefficients, as expressed by log *P* values, are to some extent, correlated to certain biological endpoints.

$$log (1/C) = k1(log P) + k2\sigma + k3$$

Conc. of compound required to produce a standard response in a given t

Logarithm of the molecule's partition coefficient (1-octanol/water)

Hammet Parameter (molecule's electronic characteristics)



'60: QSAR Golden Age



April 20, 1964

CORRELATION OF BIOLOGICAL ACTIVITY AND CHEMICAL STRUCTURE

1617

[CONTRIBUTION FROM THE DEPARTMENT OF CHEMISTRY, POMONA COLLEGE, CLAREMONT, CALIFORNIA]

 $\rho - \sigma - \pi$ Analysis. A Method for the Correlation of Biological Activity and Chemical Structure

By Corwin Hansch and Toshio Fujita1

RECEIVED AUGUST 19, 1963

Using the substituent constant, σ , and a substituent constant, π , defined as $\pi = \log P_{\rm X} - \log P_{\rm H}$ ($P_{\rm H}$ is the partition coefficient of a parent compound and $P_{\rm X}$ that of a derivative), regression analyses have been made of the effect of substituents on the biological activity of benzoic acids on mosquito larvae, phenols on gram-positive and gram-negative bacteria, phenyl ethyl phosphate insecticides on houseflies, thyroxine derivatives on rodents, diethylaminoethyl benzoates on guinea pigs, and carcinogenic compounds on mice.

Journal of Medicinal Chemistry

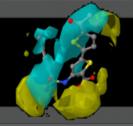
@ Copyright 1964 by the American Chemical Society

Volume 7, Number 4

July 6, 1964

A Mathematical Contribution to Structure-Activity Studies

SPENCER M. FREE, JR., AND JAMES W. WILSON



Free-Wilson



Journal of Medicinal Chemistry

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Volume 7, Number 4

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A Mathematical Contribution to Structure-Activity Studies

SPENCER M. FREE, JR., AND JAMES W. WILSON

X R N(CH₃)₂ OH OH OH OH

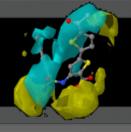
where R was H or CH_3 ; X was Br, Cl, or NO_2 ; and Y was NO_3 , NH_2 , or CH_3CONH .

BIOLOGICAL ACTIVITY OF TEN TETRACYCLINES

	Compound identification									
Com-	R		X			· · · · · · · · · · · · · · · · · · ·			logical	
pound	H	$CH_{\mathfrak{p}}$	NO_2	-C1]3 r	NO_2	$\mathrm{NH_2}$	CH_3CONH	activity	
III	1		1			}			60	
IV	1			1		1			21	
V	1				1	1			1.5	
VI	1			1			i		525	
VII	1				1		1		320	
VIII	l		1				I		275	
IX		1	1				1		160	
X		1	1					1	1.5	
XI		ł			1		1		140	
XII		i			1			1	75	

Contribution of Structural Changes^a

Side chain positions												
R		X		Ž.								
a[H]	75	b[Cl]	84	$c[{ m NH_2}]$	123							
$a[{ m CH_3})$	-112	$b\left[\mathbf{Br} ight]$	- 16	$c[\mathrm{CH_3CONH}]$	18							
		$b[\mathrm{NO}_2]$	-26	$c[NO_2]$	-218							



Mixed Hansch/Free-Wilson



Full Papers

A Combined Hansch/Free-Wilson Approach as Predictive Tool in QSAR Studies on Propafenone-Type Modulators of Multidrug Resistance^[1]

Claudia $\text{Tmej}^{a)}$, Peter Chiba $^{b)}$, Mario Huber $^{a)}$, Elisabeth Richter $^{b)}$, Manuela Hitzler $^{b)}$, Klaus-Jürgen Schaper $^{c)}$, and Gerhard Ecker $^{a)}$ *

Arch. Pharm. Pharm. Med. Chem.

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0365-6233/98/0708/0233 \$17.50 +.50/0