Medicinal Chemistry-I

DRUGS & DRUG TARGETS

Bratislava, 2014

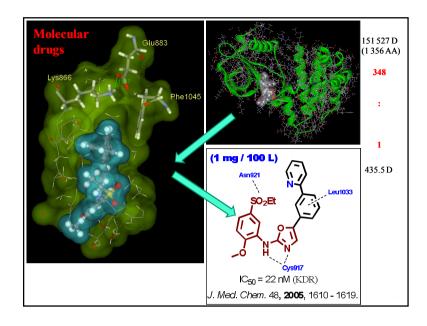
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What is medicinal chemistry?

- not a basic chemistry course for medical students
- highly interdisciplinary research dedicated to development of new drugs (not only)







What is drug?

- In medicinal chemistry, the chemists design and synthesize a pharmaceutical agent that has a desired biological effect on the human body or some other living systems.
- **Drugs** in general are **compounds** which interact with a biological system to **produce a biological response**. No one is totally safe, they vary in side effects. Dose level of a compound determines whether it will act as a **medicine or as a poison**.

(it is a dose that make from the compound a poison like: 100 aspirin tablets or 1 L bottle of whisky or 9 kg of spinach)

Chronology of Drug development

- □ selection of disease (cardiovascular, autoimmune, infectious, hereditary, mental, cancer ...)
- ☐ molecular mechanism of the pathology (medicine, molecular biology)
- □ selection of a key biomolecule to influence
- □ new active structure/compound identification: in Silico design, HTS, of organic molecules possessing appropriate drug-like properties (biologists, computer chemists)
- ☐ organic synthesis (chemists)
- ☐ biological or biophysical **assavs** (biologists)
- □ **optimization** of activity and other molecular properties (solubility, toxicity etc.)
- ☐ IP protection + clinical trials + up-scale synthesis + authority approval

What kind of compounds are drugs?

• Different inorganic, more likely organic compounds and biomolecules (proteins, antibodies, siRNA...) that activates or inhibits the function of a target with benefit to the patient







□ active

(stereoelectronicaly compatibile with target binding place)

- **possesing low toxicity** (selectivity, antitargets: e.g. cytochrome P450 enzymes, heart potassium ion channel hERG, P-glycoprotein...etc.)
- good bioavailablity (complex of physico-chemical and pharmacological properties ensuring drug-likeness)

How many new drugs reach the market vearly?

- DD is highly interdisciplinary science that is time and resources consuming process:
- 10 years / from 870 000 000 to 2 000 000 000 USD /1 new drug Adams C, Brantner V. Health Affairs (Millwood) 2006 25 420-8.
- global production ca 24 innovative drugs (new chemical entity) / year

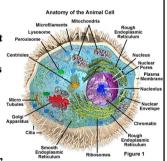
(2009: 26, 2008: 25, 2007: 18, 2006: 22, 2005: 26, 2004: 24, 2003: 26, **2002**: 28, **2001**: 23, **2000**: 26)

• Many failures have been recorded in high stages of drug development, even in clinical trials) Where is a problem?

Drug-likeness was often missing. Computer aided drug design is preferred.

A structure of a cell

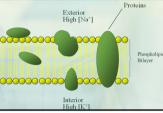
- Human, animal and plant cells are eukaryotic cells
- The nucleus contains the genetic blueprint for life (DNA)
- The fluid contents of the cell are known as the cytoplasm
- Structures within the cell are known as
- Mitochondria are the source of energy production
- Ribosomes are the cell's protein 'factories'
- Rough endoplasmic reticulum is the location for protein synthesis

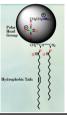


Cell membrane – protects its compartment

- CM composes from phospholipid bilayer, the hydrophobic tails interact with each other by van der Waals interactions and are hidden from the aqueous media
- The polar head groups (phosphatidylcholine) interact with water at the inner and outer surfaces of the membrane
- The cell membrane provides a **hydrophobic barrier** around the cell, **preventing the passage of water and polar molecules**. Proteins (receptors, ion channels and carrier proteins) are present, floating in the cell membrane.



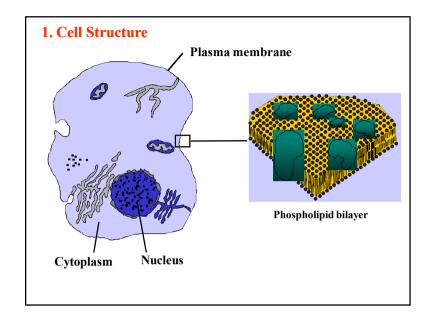




Cells in a human body

- Human body consists from up to 100 trillion (10E14) cells organized in different organs and tissues (ca 200) that operate on the molecular level (chemical reactions keeping body healthy and functional, homeostasis).
- Drug act on molecular targets in cell membrane or within the cells themselves.





Did you know? The length of all joined DNA from one adult body (111 mld km) is more as the distance between Earth and Pluto (7.5 mld km)!

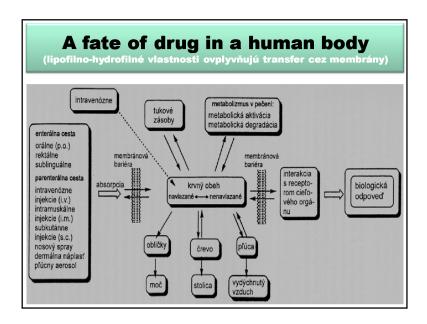
Adult human body consists from ca 3.72 x 10E13 cells. Ann Hum Biol 2013 40 471. http://www.ncbi.nlm.nih.gov/pubmed/23829164

Current lenght of human DNA is ca 3 m. http://hypertextbook.com/facts/1998/StevenChen.shtml

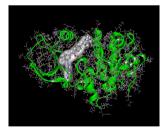
Length of all joined human DNA from one adult body is: $3.72 \times 3 \times 10E13 \text{ m} = 11.16 \times 10E10 \text{ km} = 111 \text{ mld km} !!!$

Earth Neptun distance is 4.4 mld km. http://www.universetoday.com/21628/how-far-is-neptune-from-earth/

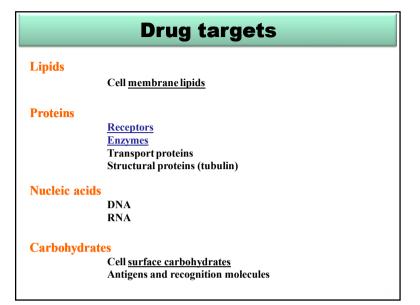
Earth Pluto distance is 7.5 mld km. http://www.universetoday.com/14313/how-far-is-pluto-from-earth/

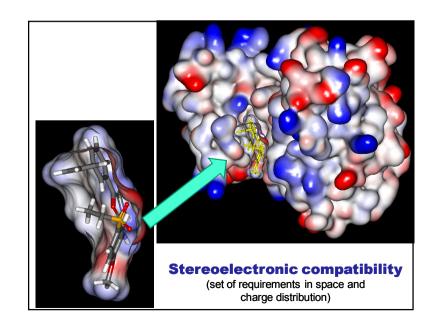


• **Drug targets are macromolecules** that have a binding site into which the drug fits and binds.



 Most drug bind to their targets by means of intermolecular bonds (electrostatics or ionic interactions, hydrogen bonds, van der Waals interactions).





Biogenic aminoacids

© COO H H R

• Unpolar (8) – (lipophilic)

Alanine	Valine	Leucine
(Ala; A)	(Val; V)	(Leu; L)
Me-	ⁱ Pr-	ⁱ Bu-
Isoleucine	Methionine	Phenylalanine
(ILE; I)	(Met; M)	(Phe; F)
*Bu-	CH ₃ S(CH ₂) ₂ -	PhCH ₂ -
Tryptophan (Try; W) (indol-3-yl)CH ₂ -	Proline (Pro; P) -(CH ₂) ₃ -	

Biogenic aminoacids

covalent bonds > ionic bonds > hydrogen bonds > van der Waals interactions



• **Ionized (5)**– (hydrophilic)

Lysine (Lys; K) H ₃ N ⁺ (CH ₂) ₄ -	Arginine (Arg; R) H ₂ N(NH ₂ +)CNH(CH ₂) ₃ -	Histidine (His; H)
Aspartic acid (Asp; D)	Glutamic acid (Glu; E) -OOC(CH ₂) ₂ -	

covalent bonds > ionic bonds > hydrogen bonds > van der Waals interactions

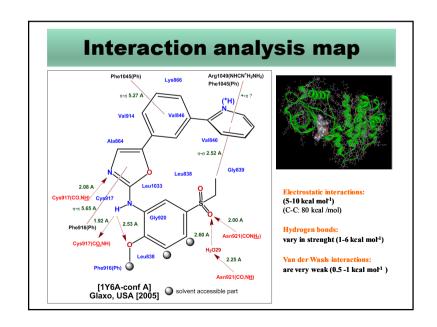
Biogenic aminoacids



• **Polar (7)** – (hydrophilic)

Glycine (Gly; G) H-	Serine (Ser; S) HOCH ₂ -	Threonine (syn; 2S,3R) (Thr; T) HOCHCH ₃
Cysteine (Cys; C) HSCH ₂ -	Tyrosine (Tyr; Y) para-HOPh-CH ₂ -	Asparagine (Asn; N) NH ₂ COCH ₂ -
Glutamine (Gln; Q) NH ₂ CO(CH ₂) ₂ -		

covalent bonds > ionic bonds > hydrogen bonds > van der Waals interactions

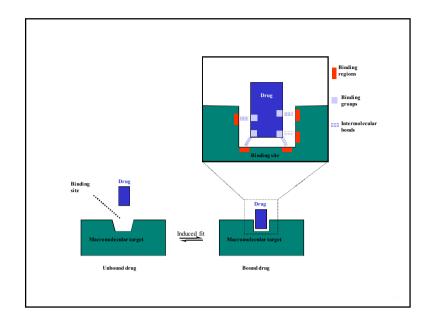


Drug / target binding terms

- Drug targets are large molecules macromolecules
- Drugs are generally much smaller than their targets
- Drugs interact with their targets by binding to binding sites
- Binding sites are typically hydrophobic hollows or clefts on the surface of macromolecules
- Binding interactions typically involve intermolecular bonds
- Most drugs are in equilibrium between being bound and unbound to their target
- Functional groups on the drug are involved in binding interactions and are called binding groups
- Specific regions within the binding site that are involved in binding interactions are called binding regions

Induced fit

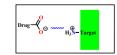
- Binding interactions usually result in an induced fit where the binding site changes shape to accommodate the drug
- The induced fit may also alter the overall shape of the drug target important to the pharmacological effect of the drug

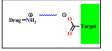


Intermolecular	bonding	forces
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Electrostatic or ionic bond

- Strongest of the intermolecular bonds
 (20-40 kJ mol⁻¹) (5 10 kcal/mol, C-C: 80 kcal /mol, C-H 110 kcal/mol)
- Takes place between groups of opposite charge
- The strength of the ionic interaction is <u>inversely proportional</u> to the <u>distance</u> between the two charged groups
- Stronger interactions occur in hydrophobic environments
- The strength of interaction <u>drops off less rapidly with distance than with other forms of intermolecular interactions</u>
- Ionic bonds are <u>the most important initial interactions</u> as a drug enters the binding site



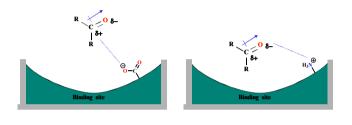


C-N

O=O 116 (2 x 58) C=O 187* (2 x 93.5) C=C 145 (2 x 72.5)

Ion-dipole interactions

- Occur where the charge on one molecule interacts with the dipole moment of another
- Stronger than a dipole-dipole interaction
- Strength of interaction falls off less rapidly with distance than for a dipole-dipole interaction



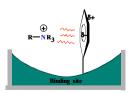
Dipole-dipole interactions S-O Dipole moment Localised dipole moment Binding site Binding site

Dipole-dipole interactions

- Can occur if the drug and the binding site have dipole moments
- Dipoles align with each other as the drug enters the binding site
- Dipole alignment orientates the molecule in the binding site
- Orientation is beneficial if other binding groups are positioned correctly with respect to the corresponding binding regions
- Orientation is detrimental if the binding groups are not positioned correctly
- The strength of the interaction decreases with distance more quickly than with electrostatic interactions, but less quickly than with van der Waals interactions

Induced dipole interactions

- occur where the <u>charge on one molecule induces a dipole on</u> another
- between a <u>quaternary ammonium ion and an aromatic ring</u> (e.g. Lys, Arg)

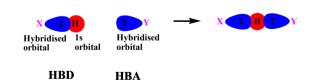


Hydrogen bonds

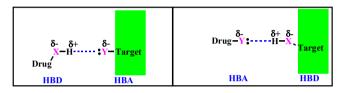
- · Vary in strength
- Weaker than electrostatic interactions but stronger than van der Waals interactions
- A hydrogen bond takes place between an electron deficient hydrogen and an electron rich heteroatom (N or O)
- The electron deficient hydrogen is usually attached to a heteroatom (O or N)



• Optimum orientation is where the X-H bond points directly to the lone pair on Y such that the angle between X, H and Y is 180°



- The electron deficient hydrogen is called a hydrogen bond donor (HBD)
- The electron rich heteroatom is called a hydrogen bond acceptor (HBA)
- **HB distance** \leq **2.5** Å (e.g. C-C bond is 1.54 Å, 0.154 nm)



Hydrogen bonds

- strong hydrogen bond acceptors (HBA)
 - carboxylate ion, phosphate ion, tertiary amine RCOO, RP(=O)(O), R₃N
- moderate hydrogen bond acceptors (HBA)
 - carboxylic acid, amide oxygen, ketone, ester, ether, alcohol RCOOH, RC(=O)NHR', RC(=O)R', RCOOR', ROH', ROH
- poor hydrogen bond acceptors (HBA)
 - sulphur, fluorine, chlorine, aromatic ring, amide nitrogen, aromatic amine

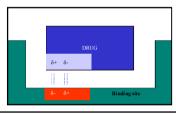
S, F, Cl, Ph, RC(=O)NHR', ArNH-

- good hydrogen bond donors (HBD)
 - quaternary ammonium ion R₃HN⁺

Van der Waals interactions

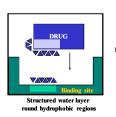
- Very weak interactions (2-4 kJ mol⁻¹)
- Occur between hydrophobic regions of the drug and the target
- Transient areas of high and low electron densities cause temporary dipoles
- Interactions drop off rapidly with distance
- Drug must be close to the binding region for interactions to occur
- The overall contribution of van der Waals interactions can be crucial to binding

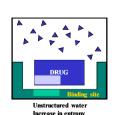




6. Hydrophobic interactions

- Hydrophobic regions of a drug and its target are not solvated
- Water molecules interact with each other and form an ordered layer next to hydrophobic regions - negative entropy
- Interactions between the hydrophobic regions of a drug and its target 'free up' the ordered water molecules
- Results in an increase in entropy that is beneficial to binding energy

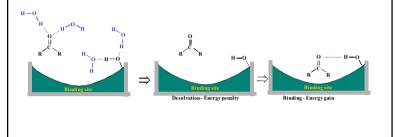






5. Desolvation penalties

- Polar regions of a drug and its target are solvated prior to interaction
- · Desolvation is necessary and requires energy
- The energy gained by drug-target interactions must be greater than the energy required for desolvation



Basic terms in medicinal chemistry





LIGAND (small organic molecule possessing target affinity, that has
to be stereoelectronically compatible with binding pocket)

- ☐ HIT an compound identified in a screen with **confirmed structure** and **activity** (need to be developed into a lead compound)
- □ LEAD active compound with convenient properties: drug-likeness, solubility, synthetic feasibility, novel structure (patentable)
- □ DRUG CANDIDATE high activity, good selectivity, low toxicity, good preclinical efficiency
- □ DRUG successfull in clinical trials, approved by FDA, EMEA for the market
- BIOAVAILABILITY basic condition to reach the target in the body
- DRUG-LIKENESS complex properties including ADME/Tox

(Absorption Distribution Metabolism Excretion / Toxicity)

Recommended literature and other sources

