Lecture 1 Introduction to pharmaceutical chemistry

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Chemistry and pharmacology

- Chemistry is the defining science of pharmacy.
- To understand anything about a drug, we need to know:
- The synthesis of the drug.
- The determination of its purity.
- The formulation into a medicine.
- The dose given, the absorption and distribution around the body.
- The molecular interaction of drug with its receptor.
- The metabolism of the drug.
- The elimination of drug from the body.
- Understanding of the chemical structure of the drug and how this chemical structure influences the properties and behavior of the drug in the body.



References used in this course are mainly:

- 1. Essentials of Pharmaceutical Chemistry by Donald Cairns, UK.
- 2. Fundamentals of Medicinal Chemistry by Gareth Thomas, UK.
- 3. Youtubes and other online lectures.

Drugs behave in solution as either weak acids or weak bases.

- Most of the drugs used in medicine are small organic molecules that behave in solution as either weak acids or weak bases.
- To understand and appreciate these compounds a study must be made of simple acid-base theory.
- in 1887, the Swedish chemist Svante August Arrhenius suggested that solutions that conduct electricity (so-called electrolytes) do so..... because they dissociate into charged species called ions.



3

Cations and anions

- Positively charged ions (or cations) migrate towards the negative terminal, or cathode.
- While negatively charged ions (or anions) migrate towards the positive terminal, or anode.
- It is this movement of ions that allows the passage of electric current through the solution.



Electrolytes and mass of action

- Compounds may be classified as strong electrolytes, which dissociate almost completely into ions in solution, or as weak electrolytes, which only dissociate to a small extent in solution.
- Since strong electrolytes are almost completely dissociated in solution, measurement of the equilibrium constant for their dissociation is very difficult.
- For weak electrolytes, however, the dissociation can be expressed by the law of mass action in terms of the equilibrium constant.
- Considering the reaction: $A+B\rightarrow C+D$:
- The equilibrium constant (K) for the reaction is given by the product of the concentrations of the reaction products divided by the product of the concentrations of the reactants, or

$$K = \frac{[C] \times [D]}{[A] \times [B]}$$

5

Equilibrium

- If the equilibrium lies to the right-hand (or products) side, the numerator in the above expression will be greater than the denominator, and K will be greater than 1.
- Conversely, if the reaction does not proceed very far and the equilibrium lies closer to the left hand side, [A] × [B] will be larger than [C] × [D] and K will be less than 1.
- The law of mass action can be applied to the dissociation of water, a weak electrolyte widely used as a solvent in biological and pharmaceutical systems: Kw = [H+] × [OH–]
- Kw is called the ionic product or autoproteolysis constant of water

 $H_2 O \rightleftharpoons H^+ + O H^-$

The equilibrium constant for this reaction is given by

$$K = \frac{[\mathrm{H}^+] \times [\mathrm{OH}^-]}{[\mathrm{H}_2\mathrm{O}]} \qquad \qquad \mathbf{p}\mathbf{H} = -\log[\mathbf{H}^+] = \log\frac{1}{[\mathbf{H}^+]}$$

<equation-block><section-header><equation-block><equation-block>Dissociation of weak acids and bases• Acids are compounds that ionise to release hydrogen ions, or protons, to their surroundings. Bases are compounds that can accept hydrogen ions. This is called the Brønsted-Lowry definition of acids and bases• The dissociation of a weak acid is usually represented as follows: HA = H++A=.• Store the most common solvent in pharmaceutical and biological systems is water, the ionization of a weak acid is better represented as: $HA + H_2 O = H_3 O^+ + A^-$ where $H_3 O^+$ is a hydroxonium ion, and the ionisation of a base can be represented as $B + H_2 O = BH^+ + OH^ K_a = \frac{(H^+) \times [A^-]}{(HA)}$ Consider any weak acid HA, which dissociates as shown below: $HA = H^+ + A^-$ The equilibrium constant for this reaction is given, as before, by $K_b = \frac{(BH^+) \times [OH^-)}{[B]}$

р	H and pOH	
The following equilibrium is extremely useful	because they allow the pH of so	lutions of weak acids and bases to be
calculated if the concentrations and dissociation	constant are known.	
For weak electrolytes, α is very small and may be neglected so $(1 - \alpha)$ is approximately equal to 1. The simplified expression may now be written as $K = \alpha^2 \alpha$	As before α is very small and can be neglected, sequal to 1.	o $(1 - \alpha)$ is approximately
where c is the concentration, in moles per litre, and α is the degree of ionisation of the acid. Then $\alpha = \sqrt{\frac{K_a}{K_a}}$	$\alpha^{2} = K_{b}/c$ $\alpha = \sqrt{(K_{b}/c)}$ From above,	
The pH of the solution can now be determined: $[H^+] = \alpha c$	$[OH^{-}] = c\alpha$ Therefore, $[OH^{-}] = c\sqrt{(K_b/c)} = \sqrt{(K_b c)}$	$\mathbf{pH} = \mathbf{p}K_{\mathbf{w}} - \frac{1}{2} \mathbf{p}K_{\mathbf{b}} + \frac{1}{2} \log c$
Therefore, $[\mathbf{H}^+] = c \sqrt{\left(\frac{K_a}{c}\right)} = \sqrt{(K_a c)}$	However, $[OH^-] = \frac{K_w}{[H^+]}$ Therefore,	
Taking logarithms, $\log[\mathbf{H}^+] = \frac{1}{2} \log K_a + \frac{1}{2} \log c$	$\frac{K_{\rm w}}{[{\rm H}^+]} = \sqrt{(K_{\rm b}c)}$	$\mathbf{pH} = \frac{1}{2} \mathbf{p}K_{\mathbf{a}} - \frac{1}{2} \log c$
Multiplying throughout by -1 gives $-\log[H^+] = -\frac{1}{2}\log K_a - \frac{1}{2}\log c$	$[\mathrm{H}^+] = \frac{K_\mathrm{w}}{\sqrt{(K_\mathrm{b}c)}}$	
Therefore, $\mathbf{pH} = \frac{1}{2} \mathbf{pK}_{a} - \frac{1}{2} \log c \qquad (1.2)$	Taking logarithms, $\log[H^+] = \log K_w - \frac{1}{2} \log K_b - \frac{1}{2} \log c$	

pKa and pKb

• It is often useful and convenient to express the strengths of acids and bases using the same term, p*K*a, and this can be done by considering the equilibria that exist between an acid and its conjugate base.

- A weak acid (HA) and its conjugate base (A-) are related as follows:
- pKa = log Ka

Consider the two carboxylic acids below: Acetic acid, CH₃COOH, $pK_a = 4.7$ Chloroacetic acid, ClCH₂COOH, $pK_a = 2.7$ $HA \rightleftharpoons H^{+} + A^{-}$ $A^{-} + H_2O \rightleftharpoons HA + OH^{-}$

From the equations above,

 $K_{\rm a} = \frac{[\rm H^+] \times [\rm A^-]}{[\rm HA]}$

and

$$K_{\rm b} = \frac{[\rm HA] \times [\rm OH^{-}]}{[\rm A^{-}]}$$

Then

$$K_{a} \times K_{b} = \frac{[\mathrm{H}^{+}] \times [\mathrm{A}^{-}]}{[\mathrm{H}\mathrm{A}]} \times \frac{[\mathrm{H}\mathrm{A}] \times [\mathrm{O}\mathrm{H}^{-}]}{[\mathrm{A}^{-}]}$$

Cancelling similar terms gives

$$K_{\rm a} \times K_{\rm b} = [{\rm H}^+] \times [{\rm OH}^-]$$

which can be rewritten as

$$K_{\rm a} \times K_{\rm b} = K_{\rm w} = 1 \times 10^{-14}$$

9

Hydolysis of salts

- When a salt is dissolved in water, the compound dissociates completely to give solvated anions and cations. This breaking of bonds by the action of water is called hydrolysis and the salt is said to be hydrolyzed.
- The pH of the resulting solution depends on whether the salt was formed from reaction of strong or weak acids and bases and there are four possible combinations.

These relationships can be summarised as follows:

Strong $acid + Strong \ base \rightarrow Neutral \ salt$ Strong $acid + Weak \ base \rightarrow Acidic \ salt$ Weak $acid + Strong \ base \rightarrow Basic \ salt$ Weak $acid + Weak \ base \rightarrow Neutral \ salt$

Amphiprotic salts

- The reactions of salts in water become more complicated if the salt in question is *amphiprotic*; that is, it can function both as an acid and a base.
- Examples of amphiprotic anions are bicarbonate (sometimes called hydrogencarbonate), HCO3-, and bisulfite (or hydrogensulfite), HSO3-. These species can donate or accept hydrogen ions in solution.

The pH of a solution of an amphiprotic salt (e.g. sodium bicarbonate, $Na^+HCO_3^-$) is given by the equation

$$pH = \frac{1}{2}(pK_{a1} + pK_{a2}) \tag{1.6}$$

11

Buffer solutions

- A buffer solution is a solution that resists changes in pH.
- If acid is added then, within reason, the pH does not fall; if base is added, the pH does not rise.
- Buffers are usually composed of a mixture of weak acids or weak bases and their salts and function best at a pH equal to the pKa of the acid or base involved in the buffer.
- The equation that predicts the behaviour of buffers is known as the Henderson-Hasselbalch equation.
- It is derived as follows, by considering a weak acid that ionises in solution:

$$\mathbf{pH} = \mathbf{pK_a} + \log \frac{[\mathbf{SALT}]}{[\mathbf{ACID}]}$$

Buffer capacity

- Buffer solutions work best at controlling pH at pH values roughly equal to the pKa of the component acid or base: that is, when the [SALT] is equal to the [ACID].
- This can be shown by calculating the ability of the buffer to resist changes in pH, which is the buffer capacity.
- *The buffer capacity* is defined as the number of moles per litre of strong monobasic acid or base required to produce an increase or decrease of one pH unit in the solution.
- When the concentrations of salt and acid are equal, the log term in the Henderson– Hasselbalch equation becomes the logarithm of 1, which equals 0.

13

Adjust the pH of a buffer

• To move the pH of the buffer solution by one unit of pH will require the Henderson-Hasselbalch equation to become:

$$pH = pK_a + \log\frac{10}{1}$$

• It will require addition of more acid or base to move the pH by one unit from the point where pH = pKa than at any other given value of the ratio. This can be neatly illustrated by the following example.

An example of a buffer

Suppose 1 litre of buffer consists of 0.1 M CH₃COOH and 0.1 M CH₃COO⁻Na⁺: the pH of this buffer solution will be 4.7 (since the log term in the Henderson–Hasselbalch equation cancels). Now, if 10 mL of 1 M NaOH is added to this buffer, what will be the new pH?

Clearly, the 10 mL of NaOH will ionise completely (strong alkali) and some of the 0.1 M acetic acid will have to convert to acetate anion to compensate. The new pH will be

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pH = pK_a + \log \frac{[SALT]}{[ACID]}
pH = 4.7 + \log \frac{(0.1 + 0.01)}{(0.1 - 0.01)}
pH = 4.7 + \log \frac{0.11}{0.09}
pH = 4.79
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15

Biological buffers

- The human body contains many buffer systems, which control the pH of body compartments and fluids very effectively.
- Blood plasma is maintained at a pH of 7.4 by the action of three main buffer systems:
- 1. dissolved carbon dioxide, which gives carbonic acid (H2CO3) in solution, and its sodium salt (usually sodium bicarbonate, NaHCO3), which is responsible for most of the buffering capacity.
- 2. The other two buffers are dihydrogen phosphate (H2PO4–), also with its sodium salt.
- 3. The protein macromolecules "Proteins are polymers composed of repeating units called amino acids" which containing NH2 and COOH groups:

Proteins capable of acting as both acids and bases

- Proteins are composed of about 20 different amino acid residues, which are connected to each other by peptide bonds formed between one amino acid and its neighbor.
- The side-chain of the amino acid may be acidic (as in the case of glutamic and aspartic acids), basic (as in the case of arginine and lysine) or neutral (as in alanine).
- A protein, which may be composed of hundreds of amino acid residues, is, therefore, a polyelectrolyte whose properties depend on the balance of acidic and basic groups on the side-chains.

Proteins capable of acting as both acids and bases

- Generally, most proteins act as weak acids and form buffers with their sodium salts.
- free amino acids usually do not exist in the molecular form, but instead both the amino and carboxyl groups ionise to form an internal salt.
- These internal salts are known by the German word zwitterion ('dipolar ion'), and formation of the zwitterion makes the amino acid very polar and, therefore, very soluble in v





19

pKa of the drug

• Equation (1.8) applies to drugs that are weak acids and allows the fraction of the total dose that is ionised to be calculated for any pH if the pKa of the drug is known.

• The equation is sometimes written as the percentage ionised, which is simply given by

Fraction ionised =
$$\frac{1}{1 + \operatorname{antilog}(pK_a - pH)}$$
 (1.8)

pKa values of drug molecules

- Most compounds used in medicine are either weak acids or weak bases. This means that the range of possible pKa values encountered in drug molecules is huge.
- It is important to remember that the value of the pKa for a drug tells you absolutely nothing about whether the compound is an acid or base.
- Many weak bases have pKa values of 2 to 4. Similarly, while a basic drug like cocaine has a pKa of 9.5, this does not mean that all compounds with a pKa greater than 7 are bases.
- Only a thorough understanding of chemical structure and a knowledge of the functional groups that confer acidity or basicity on a molecule will allow the correct prediction of the acidic or basic nature of a molecule.

pKa of some common drugs			
Table 1.1 pKa values of some common drug	S		
Drug	рК _а value		
Acidic drugs			
Aspirin	3.5		
Paracetamol	9.5		
Phenobarbital	7.4 (first ionisation)		
Basic drugs			
Cocaine	8.6		
Diazepam	3.3		
Diphenhydramine	9.0		
Amphoteric drugs			
Morphine	8.0 (amine), 9.9 (phenol)		
Adrenaline (epinephrine)	8.7 (amine), 10.2, 12.0 (phenols)		

Lecture 2 Partition coefficient *P* and Biopharmacy

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1

Partition coefficient and biopharmacy

- A drug has to pass through a number of biological membranes in order to reach its site of action.
- Consequently, organic medium/aqueous system partition coefficients were the clear parameters to use as a measure of the ease of movement of the drug through these membranes.
- The accuracy of the correlation of drug activity with partition coefficients will depend on the solvent system used as a model for the membrane.
- What is partition coefficients ?





Partition coefficient and biopharmacy

- When a substance (or *solute*) is added to a pair of immiscible solvents, it distributes itself between the two solvents according to its affinity for each phase.
- A polar compound (e.g. a sugar, amino acid or ionized drug) will tend to favour the aqueous or polar phase.
- whereas a non-polar compound (e.g. an unionized drug) will favour the nonaqueous or organic phase.
- The added substance distributes itself between the two immiscible solvents according to the partition law, which will '' divide" itself between two immiscible solvents in a constant ratio of concentrations.



Partition coefficient and biopharmacy

 $P = \frac{[\text{organic}]}{[\text{aqueous}]}$

An example

of partition

coefficient

- This constant ratio is called the "*partition coefficient*" of the substance, and may be expressed mathematically as below:
 - Where *P* is the partition coefficient of the substance; [organic] is the concentration of substance in the organic, or oil phase; and [aqueous] is the concentration of substance in the water phase.

where P is the partition coefficient of the substance; [organic] is the concentration of substance in the organic, or oil phase; and [aqueous] is the concentration of substance in the water phase.

As an example, consider the distribution of 100 mg of a drug between 50 mL of an organic solvent (e.g. ether, chloroform or octanol) and 50 mL of water. The drug is added to the two immiscible solvents in a separating funnel and allowed to equilibrate. When the organic layer is analysed, it is found to contain 66.7 mg of compound. From these data the partition coefficient and the percentage of the drug extracted into the organic layer can be calculated (see Fig. 2.1).

The mass of drug in the water phase = 100 - 66.7 mg = 33.3 mg; the concentration of drug in the organic phase = $66.7/50 = 1.33 \text{ mg mL}^{-1}$, and the concentration of drug in the water phase = $33.3/50 = 0.67 \text{ mg mL}^{-1}$. Therefore, the partition coefficient is given by

$$\frac{[\text{organic}]}{[\text{aqueous}]} = \frac{1.33 \text{ mg mL}^{-1}}{0.67 \text{ mg mL}^{-1}} = 2$$





• Part of medicinal chemistry, the science of rational drug design, *involves structureactivity relationships*, where the partition coefficient is used in mathematical equations that try to relate the biological activity of a drug to its physical and chemical characteristics.

QSAR and physicochemical properties of Drugs



- QSAR stands for Quantitative Structure-Activity Relationship.
- It is a computational modeling technique used in chemistry and pharmacology to predict the biological activity, potency, or other properties of chemical compounds based on their structural characteristics.
- QSAR models relate the chemical structure of a molecule to its biological or physicochemical properties through mathematical equations.

9

Ionization and pH

- This relationship above only applies if the solute in question does not ionize at the pH of measurement.
- If the solute is a weak acid or weak base (and a huge number of drugs are), then ionization to form an anion or a cation will considerably alter the solubility profile of the drug.
- A fully ionized species will be much more soluble in water than the unionized acid or base, and so the above ratio will vary depending on the pH at which the measurement was carried out.



P value and ionazation

- There are two ways round this problem: the experimental conditions are adjusted to ensure that the measured *P* is the partition coefficient of the unionized molecule (this means that the *P* value for acids is measured at low pH when the acid is unionized.
- OR better, the ratio above is redefined as the *apparent partition coefficient*.
- It depends upon the pH of the solution.

$P_{\rm app} = P \times f_{\rm unionised}$

where $f_{\text{unionised}}$ equals the fraction of the total amount of drug union that pH. It follows that if $f_{\text{unionised}} = 1$ then $P_{\text{app}} = P_{\text{true}}$ and the com is unionised.

To illustrate the effect of ionisation, consider again the drug example above. If the pH of the aqueous phase is adjusted so that th becomes 66.7% ionised, only 40 mg of the drug partitions into the c phase (since the ionised drug will be less soluble in the organic solven the partition coefficient can be recalculated (see Fig. 2.2).



P value and quantitative structure-activity relation- ships (QSAR)

- The range of possible values of P found in drug molecules is huge, from small fractions through to values of several thousand.
- For this reason, it is common to quote the logarithm (to the base 10) of the partition coefficient, or log *P*.
- This is particularly true in *quantitative structure-activity relation- ships* (QSAR), where the physicochemical properties of a drug (such as hydrophobicity, steric interactions or electronic effects) are quantified and an equation is derived that can be used to predict the biological activity of other, similar drugs.





- These parameters are used to represent properties such as lipophilicity, shape and electron distribution, which are believed to have a major influence on the drug's activity.
- The main properties of a drug that appear to influence its activity are its, lipophilicity, the electronic effects within the molecule and the size and shape of the molecule (steric effects).

Lipophilicity



- Lipophilicity is a measure of a drug's solubility in lipid membranes.
- This is usually an important factor in determining how easily a drug passes through lipid membranes
- The electronic effects of the groups within the molecule will affect its electron distribution, which in turn has a direct bearing on how easily and permanently the molecule binds to its target molecule.
- Drug size and shape will determine whether the drug molecule is able to get close enough to its target site in order to bind to that site.
- Two parameters are commonly used to represent lipophilicity, namely the partition coefficient (P) and the lipophilicity substituent constant (p). The former parameter refers to the whole molecule whilst the latter is related to substituent groups.



Electronic effects

- The distribution of the electrons in a drug molecule has a considerable influence on the distribution and activity of a drug.
- In general, nonpolar and polar drugs in their unionized form are more readily transported through membranes than polar drugs and drugs in their ionized forms.
- Furthermore, once the drug reaches its target site the distribution of electrons in its structure will control the type of bond it forms with that target, which in turn affects its biological activity

17

Steric effects

- The first parameter used to show the relationship between the shape and size (bulk) of a drug, the dimensions of its target site and the drug's activity was the Taft steric parameter (Es).
- It was followed by Charton's steric parameter (n), Verloop's steric parameters and the molar refractivity (MR) amongst others.
- The most used of these additional parameters is probably the molar refractivity.

$$E_{\rm s} = \log \frac{k_{\rm (XCH_2COOCH_3)}}{k_{\rm (CH_3COOCH_3)}} = \log k_{\rm (XCH_2COOCH_3)} - \log k_{\rm (CH_3COOCH_3)}$$

Table 4.6	Examples	of the Taft	steric parameter	E
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Group	$E_{\rm s}$	Group	$E_{\rm s}$	Group	$E_{\rm s}$
H–	1.24	F-	0.78	CH ₃ O-	0.69
CH ₃ -	0.00	Cl-	0.27	CH ₃ S-	0.19
C_2H_5-	-0.07	F ₃ C-	-1.16	PhCH ₂ -	-0.38
(CH ₃) ₂ CH-	-0.47	Cl ₃ C-	-2.06	PhOCH-	-0.33



Molar refractivity (MR)

• The molar refractivity is a measure of both the volume of a compound and how easily it is polarized. It is defined as:

$$\mathrm{MR} = \frac{(n^2 - 1)M}{(n^2 + 2)\rho}$$

• where n is the refractive index, M the relative mass and r the density of the compound. The M/r term is a measure of the molar volume whilst the refractive index term is a measure of the polarizability of the compound.

 Table 4.7
 Examples of calculated MR values. Reproduced by permission of John Wiley and Sons

 Ltd. from Hansch C. and Leo A.J. Substituents Constants for Correlation Analysis in Chemistry and

 Biology (1979)

Group	MR	Group	MR	Group	MR
H–	1.03	F–	0.92	CH ₃ O-	7.87
CH ₃ -	5.65	Cl-	6.03	HO-	2.85
C2H5-	10.30	F ₃ C-	5.02	CH ₃ CONH-	14.93
(CH ₃) ₂ CH-	14.96	O ₂ N-	7.63	CH ₃ CO-	11.18

Lecture 3 Measurement of partition coefficient

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1

Experimental measurement to determine the partition coefficient: Shake flask method

- These are the original *shake flask method*, the use of *thin-layer chromatography* or the use of *reversed-phase*, *high-performance liquid chromatography*.
- Shake flask method:
- In the shake flask method, the drug whose P is to be determined is traditionally added to a separating funnel containing the two immiscible phases, although it works just as well to use a centrifuge tube (and requires less sample).
- The two immiscible phases chosen are usually 1-octanol and pH 7.4 buffer.
- Octanol is used in partition coefficient work because the answers obtained from octanol seem to correlate best with biological data obtained in vivo.



Continue with...Shake flask method

- This may be because the eight carbon atoms are essentially hydrophobic (or water-hating) and the one hydroxyl group is hydrophilic (water-loving) and together they give the closest balance to that found in human cell membranes.
- The aqueous buffer at pH 7.4 represents aqueous compartments within the body, e.g. blood plasma.
- The two phases are thoroughly mixed to give buffer-saturated octanol in the top phase and octanolsaturated buffer in the bottom. .
- The concentration in the octanol phase is found by subtraction and the value of P is calculated.
- This method works perfectly well if there is sufficient sample and the drug possesses a chromophore to allow spectroscopic assay of the aqueous phase.
- Five extractions of 10 mL organic phase will remove more compound than one extraction of 50 mL, even though the total volume of organic solvent used is the same.

Experimental measurement to determine the partition coefficient: Thin-layer chromatography (TLC)

- In this technique, the *R*f value of the drug is related mathematically to the partition coefficient.
- A thin-layer plate, or a paper sheet, is pre-coated with organic phase (usually paraffin or octanol) and allowed to dry.
- Sample is applied to the origin and the plate is allowed to develop.
- The mobile phase used is either water or a mixture of water and a miscible organic solvent (such as acetone) to improve the solubility of the drug.
- Once the plate has developed, the spots are visualized (using an ultra- violet lamp if the drug possesses a chromophore, or iodine vapor if it does not) and the *R*f for each spot is determined.



³

*R*f is the distance and *k* is a constant for the given system

$$P=\frac{k}{(1/R_{\rm f})-1}$$

- The *R*f is the distance moved by the spot divided by the distance moved by the solvent .
- The *R*f can be related to the partition coefficient by equations of the type
- where k is a constant for the given system, which is determined by running a number of standard compounds of known P in the system and calculating k.
- The TLC method of determining *P* works best for compounds of similar structure and physical properties.
- The advantages of using this technique to determine *P* are that many compounds can be run simultaneously on one plate, and very little sample is required.

High-performance liquid chromatography (HPLC)

- This method of analysis relies on the same chemical principles as the determination by TLC.
- *The retention time*, as its name suggests, is the time taken for the sample to elute from the HPLC column.
- The major drawback with using this technique to determine *P* is detecting the drug if it does not possess a chromophore, when a UV detector cannot be used.
- There are some advantages to the HPLC method of determining *P*, namely that HPLC does not require much sample and that the sample does not have to be 100% pure.



Drug absorption, distribution and bioavailability

- The study of the fate of a drug administered to an organism is called *pharmacokinetics*.
- This discipline involves measuring or predicting the *absorption*, *distribution*, *metabolism and excretion* (usually known by the acronym ADME) of the drug in the body.
- Pharmacokinetics is an opposed to *pharmacodynamics*, which is the study of mechanisms of drug action.





Drug absorption

- The drug is not considered to be *in* the body until it has been absorbed across the gut wall and into the bloodstream by absorption across a biological membrane.
- The case of drugs acting on the brain or spinal cord (the central nervous system, or CNS) the drug must partition across the *blood-brain barrier* to gain access to the CNS.
- The CNS does require low-molecular-weight molecules to grow and function.



The cell membrane and absorption

- A cell membrane is composed of a bilayer of fatty molecules known phospholipids.
- These compounds are amphoteric in nature, possessing a non-pol region of hydrocarbon chains that are buried inside the cell membrar and a polar region comprising negatively charged phosphoric acid hegroups.
- Buried within this lipid bilayer are large globular protein molecule These macromolecules function as ion channels (e.g. the Na+ channel nerve membranes), transmembrane receptors (like the β adrenoceptc or transport proteins



Roles of fats in the cell membrane

- In cell membranes, cholesterol increases membrane rigidity and is essential for maintaining the integrity of the membrane without cholesterol your cells would leak.
- The lipid bilayer of the cell membrane presents a significant barrier to drug transport and for a small drug molecule to travel across membranes, one of two things must happen:
- 1. The drug must cross the membrane by passive diffusion.
- 2. The drug has to be transported across the membrane, against the concentration gradient, a process called active transport.



Figure 2.4. The structures of cholesterol and phospholipids. R^1 and R^2 = palmityl, stearyl or oleyl. R^3 = ethanolamine, choline, serine, inositol or glycerol.





The pH partition hypothesis

- · Biological membranes are, essentially, non-polar or hydrophobic, due to the long hydrocarbon chains of the phospholipid molecules.
- For a drug to cross a membrane of this type, the drug must pass from the aqueous solution of the extracellular fluid, through the lipid membrane to the aqueous solution of the intracellular fluid., i.e. the drug must be sufficiently soluble in both the aqueous and the lipid phases to succeed.



ogical membrane on the steady-state distribution kly acidic drug with pKa = 6



Active transport mechanisms

- Glucose and ions such as sodium and chloride must cross membranes efficiently, but they are too polar to diffuse across a phospholipid bilayer passively.
- Their transport is 'facilitated' by proteins that span the membrane and allow these chemicals to enter cells.
- If the transport occurs down a concentration gradient, the process is described as *facilitated diffusion* and does not usually require expenditure of energy in the form of hydrolysis of ATP (adenosine triphosphate).
- The protein complex diffuses across the cell membrane.





The action of local anesthetics drugs (LA)



- *Local anesthetics* are basic drugs, all derived originally from cocaine, an alkaloid obtained from the leaves of *coca*.
- Cocaine is a very effective local anesthetic, but due to a profound stimulant action on the CNS it has been replaced in most routine procedures with synthetic, non-addictive, analogues such as lidocaine etc.
- These drugs are aliphatic amines, with pKa values for their conjugate acids of approximately 8–9.

17

How does anesthetics work on nerve

- Applying the 'rule of thumb' shows that local anesthetics will exist approximately 99% ionised at blood pH (7.4).
- The site of action of most local anesthetics is a Na+ ion channel found in the cell membrane of nerve cells (or *neurons*).
- This sodium channel suggests allows Na+ ions to travel through the cell membrane to depolarise the resting membrane potential and allow the nerve cell to fire.
- Local anaesthetics block nerve conduction by attaching to the protein of the sodium channel and disrupting the flow of Na+ ions.



Excretion and reabsorption of drugs

- The same types of physicochemical processes occur when drugs are reabsorbed into the bloodstream following excretion by the kidneys.
- The two kidneys are situated at the back of the abdomen on either side of the vertebral column.
- They carry out many functions in the body, the most important of which is the production of urine and the excretion from the body of low-molecular-weight (relative molecular mass less than 68 000 daltons) water-soluble compounds, including many drugs.



19

Food and drink in Pedicel pharmacological research Calyx -· Spicy foods such as curries and chilies and flavorings such as paprika derive their hot pungent taste from the compound Placenta capsaicin Seeds -Capsaicin is found in the fruits of various species of Capsicum and is a powerful irritant causing intense pain if administered in Capsaicin a pure form. Glands Capsaicin is a non-polar compound possessing few polar groups to hydrogen-bond to water. Endocarp · This means that capsaicin is virtually insoluble in water. Mesocarp-• This is important information for people who eat spicy food. Exocarp · Drinking water with spicy is not useful because capsaicin is not soluble in aqueous solution. (you can eat yogurt with spicy to cool Apex down!).



Lecture 4 Physicochemical properties of drugs

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What do Physical and chemical Properties tells you about drugs?

- Physical property of drugs is responsible of for its actions.
- Chemical property of drugs describe the reaction of drugs extracellularly according to simple reactions like neutralization, chelation oxidation.



2

Physicochemical properties of drugs

- In this lecture we will explore the reasons why drugs behave as acids or bases and what effects ionisation has on the properties of the drug,
- The most important thing to realise about acidic and basic drugs is that values of pKa and pKb.
- It tells you absolutely nothing about whether the drug in question is an acid or a base.
- The pKa and pKb values give information about the strength of acids and bases.



What do pKa and pKb values tell you?

- The pKa and pKb values tell you the pH at which 50% of the drug is ionised, but *they do not tell you whether a drug behaves as an acid or a base in solution.*
- Amines, for example, are basic and have pKa values of approximately 9, while phenols are acidic and typically have pKa values of around 10.
- The only sure way to know whether a drug is acidic or basic *is to* learn the functional groups that confer acidity and basicity on a molecule.



Carboxylic acids R-COOH

- According to the Brønsted–Lowry definition, an acid is a substance that ionises to donate protons to its surroundings.
- In aqueous solution this is represented as $HA + H_2O \Rightarrow H_3O^+ + A^-$
- The most commonly occurring functional group conferring acidity on drug molecules is the carboxyl group, which ionises as shown below:
- A double bond in C=O is much shorter than a C-O single bond (due to sideways repulsion of the electrons in the bond) which are mesured by by X-ray diffraction












Barbiturates is an anticonvulsants or antiepileptics

- Barbiturates are cyclic imides used as hypnotics and (in the case of phenobarbital) as anticonvulsants.
- They are all derivatives of barbituric acid (which is not pharmacologically active) and differ only in their substituents on the 5-position of the ring.
- Despite Barbiturates contain nitrogen atoms, so barbiturates are not basic.
- Instead, they behave as weak acids in solution (diprotic actually, though the second ionisation is very weak).



Phenytoin

- Phenytoin is an anticonvulsant widely used in the treatment of epilepsy.
- The properties of phenytoin resemble those of barbiturates.
- It is a cyclic imide with a pKa of 8.3.
- The anion is stabilised by resonance of the negative charge onto the oxygens of the carbonyl group and the drug is usually administered as the sodium salt to increase water solubility.
- Phenytoin and barbiturates display tautomerism of the imine-imide type.







• An amine in aqueous solution will react with water to release hydroxide ions (OH-):

$$R_3N + H_2O \Rightarrow R_3NH^+ + OH^-$$

- The basicity of amines gives the availability of the lone pair of electrons on the nitrogen atom.
- If the lone pair is involved in interactions elsewhere in the molecule, then the amine will not be basic.
- Conversely, the lone pair of electrons on the amino group attached to the benzene ring is less available for reaction with protons due to delocalisation into the ring.
- This delocalisation increases the electron density of the *ortho-* and *para-*carbon atoms and means that the Ar-NH2 group does not ionise at the pH of blood.







Lecture 5 Stereochemistry of drugs structure

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<section-header> Geometry and isomers of compounds Here, we study the unusual geometry that arises around a carbon atom with four different substituents attached to it – an *asymmetric carbon atom*. The study of the three-dimensional shape of molecules is important to for drug design. Chemical compounds that have the same molecular formula, but different structural formulas are said to be isomers of each other. These structural isomers differ in their bonding sequence, i.e. their atoms are connected to each other in different ways. Stereoisomers have the same bonding sequence, but they differ in the orientation of their atoms in space.















Enantiomer and biological activity

- The common synthesis of adrenaline (epinephrine hormone), yields a racemic mixture, which has precisely 50% of the biological activity of the natural hormone.
- Once the racemate is resolved into the two pure enantiomers.
- The (*R*)-(–)-adrenaline is found to be identical to the natural hormone produced by the adrenal medulla, while the other enantiomer.
- The (S)-(+) isomer, has little or no biological activity.









• An example of a Fischer projection of lactic acid, the acid produced when milk turns sour.



The bioactivity of penicillin anti-biotics

- The mode of action of penicillin anti- biotics depends on the opposite stereochemistry of bacterial amino acids.
- In penicillin-sensitive bacteria, the organism synthesizes a cell wall to contain the high osmotic pressure produced inside the bacterial cell.
- The bacterial cell wall consists of a polysaccharide (called *peptidoglycan*), which is re-inforced by structural cross-linking of chains of polypeptide.
- The final step of the cross-linking is achieved by attaching the terminal amino acid of the cell wall, a glycine, to a D-alanine residue on an adjacent peptide chain.
- This cross-linking is catalysed by an enzyme called *transpeptidase* (or transaminase).
- Penicillins can inhibit the enzyme transpeptidase and prevent the formation of structural cross-links in the bacterial cell wall.
- The cell is weakened, becomes unable to contain the high internal osmotic pressure and bursts.

15

Mode of action of β -lactam antibiotics

- The cephalosporins which are antibiotics similar to penicillin, are known collectively as β -lactam antibiotics.
- The β-lactam ring is the 4-membered cyclic amide ring common to both classes of antibiotics and fundamental to the molecular mode of action of the drugs.
- The β-lactam ring is under immense strain and opens easily if attacked by a nucleophile.
- This is because amides contain sp2 hybridised carbon atoms, which normally have a bond angle of 120°. The bond angle of the amide in a β-lactam ring approaches 90°.
- A serine residue present in the active site of transpeptidase can attack the β-lactam ring, using the lone pair of electrons on the –OH, open the ring and so acylate the active site of the enzyme and prevent cell wall cross-linking.



Penicillin is non-toxic to humans

- Unfortunately, other nucleophiles can open a βlactam ring and inactivate penicillin.
- Some bacteria have evolved mechanisms to overcome penicillins and are said to be resistant to the drug.
- Penicillins are non-toxic to humans because we possess L-alanine, the amino acid with the opposite stereochemistry, in our proteins.
- This is an example of an important concept in drug design called *selective toxicity*, which arises when a drug is poisonous to one type of organism or cell (a bacterium in this case) but harmless to another (human cells).

17



R and **S** configurations

- The absolute configuration of atoms around a chiral centre may be drawn accurately by use of a Fischer projection and may be described (particularly in biochemistry for chiral carbohydrates and amino acids) by the D/L convention.
- This system assigns each chiral centre in a molecule a letter (*R* or *S*) and is the method of choice when assigning the configuration of chiral centres of drug molecules.
- for example: If two groups cannot be distinguished on the basis of atomic number, the next atom of the group attached to the chiral centre is considered, and so on until the priorities are clear.





Lecture 6 Drug metabolism

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Biotransformation Phase 1 reactions

		-		
Phase I reaction Upophilic Drug R Drug R AH Drug R AH Drug R AH Liver Billary syst Gastrointestinal (feces)	Phase II reaction Drug-R-CSH Drug	Kidneys	stemic culation Skin Sweat	Breats

- Definition: modification of the drug molecule via oxidation, reduction, or hydrolysis.
- Phase 1 reactions are reactions in which a new functional group is introduced into the molecule, or an existing group is converted into another (usually more water-soluble) derivative.
- Microsomal metabolism: Cytochrome P450 isozymes.
- Non-microsomal metabolism such as Hydrolysis, Monoamine oxidases and Alcohol metabolism





- Most small drug molecules are relatively lipophilic and in the body.
- Often these drugs (metabolites) are more hydrophilic than their parent drug.
- Metabolites may possess a different level of activity as the starting compound or none at all..
- In some cases, a parent drug may be inactive but is then converted into the active metabolite in the body.







- The drug metabolism involve simple chemical reactions such as oxidation (the most common), reduction and dealkylation and are influenced by a number of factors including:
- 1. Genetic factors: The science of pharmaco-genomics.
- 2. Physiological factors. These include age of the patient, gender, pregnancy and nutritional status.
- 3. Pharmacodynamic factors.
- 4. Environmental factors.

The most important and most extensively studied drug metabolism system in the body is the superfamily of cytochrome P450 monooxygenases (CYP450).



• CYP450 acts as an active electron transport system which is responsible for the oxidative metabolism of a large number of drugs and other xenobiotics such as bile acids, prostaglandins and vitamins.





Enzyme induction and inhibition

Enzyme Inhibitor & Inducer

Enzyme Inhibitor	Enzyme Inducer			
Cimetidine	Rifampicin			
Ketoconazole	Carbamazepine			
Fluconazole	Phenobarbital			

- A xenobiotic can lead to an increased rate of metabolism of a wide variety of compounds. This process is known as *enzyme induction* and is dose- dependent.
- The CYP450 enzyme system is responsible for a large number of biotransformation, so the possibility of drug interactions is very large.
- Environmental chemicals such as polycyclic aromatic hydrocarbons (PAHs) present in cigarette smoke, xanthines and flavones in foods, and other can all change the activity of CYP450 enzymes.

11

Type of inhibitions of CYP450

- *Reversible inhibitors*, such as cimetidine, which interact with the complexed iron at the active site of the enzyme to inhibit oxidation of other drugs.
- *Metabolite intermediate complexation of CYP450*. Examples of this type of inhibition include alkylamine drugs that undergo oxidation to o-nitrosoalkane derivatives.
- Mechanism-based inactivation of CYP450 (or suicide inhibition) occurs when a non-toxic drug is metabolised by CYP450 irreversibly. Examples include the antibiotic chloramphenicol and the anticancer agent cyclophosphamide.













Lecture 7 Pharmacodynamic of drugs and medicines

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- 1. When two drugs interact with the same receptor (same pharmacologic mechanism), the D-R curves will have parallel slopes.
- Drugs A and B have the same mechanism; drugs X and Y do not.
- 2. affinity can be compared only when two drugs bind to the same receptor.
- Drug A has a greater affinity than drug B.
- 3. In terms of potency, drug A has greater potency than drug B, and X is more potent than Y.
- 4. In terms of efficacy, drugs A and B are equivalent. Drug X has greater efficacy than drug Y.

Full and Partial Agonists

- Full agonists produce a maximal response-they have maximal efficacy.
- Partial agonists are incapable of eliciting a maximal response and are less effective than full agonists.
- In Figure 1-2-2, drug B is a full agonist, and drugs A and C are partial.
- Drug A is more potent than drug C , and drug B is more potent than drug C.
- At low responses, A is more potent than B, but at high responses, the reverse is true.



2. Efficacy and Potency of Full and Partial Agonists



Toxicity and the Therapeutic Index (Tl)

- Comparisons between ED50 and TD50 values permit evaluation of the relative safety of a drug (the herapeutic index), as would comparison between ED50 and the lethal median dose (LD50) if the latter is known.
- As shown in Figure I-2-5, these D-R curves can also be used to show the relationship between dose and toxic effects of a drug. The median toxic dose of a drug (TD50) is the dose that causes toxicity in 50% of a population.
- From the data shown, TI = 10/2 = 5
- Such indices are of most value when toxicity represents an extension of the pharmacologic actions of a drug. They do not predict idiosyncratic reactions or drug hypersensitivity.



Signaling Mechanisms: Types Of Drug Responsive Signaling Mechanisms

- Binding of an agonist drug to its receptor activates an effector or signaling mechanism.
- Several different types of drug-responsive signaling mechanisms are known.

• 1. Intracellular Receptors

- These include receptors for steroid, such as binding of hormones or drugs to such receptors releases regulatory proteins.
- This interaction leads to changes in gene expression.
- For example, drugs interacting with glucocorticoid receptors lead to gene expression of proteins that inhibit the production of inflammatory mediators.



ligan

GPCF

2. Membrane Receptors Directly Coupled to Ion Channels

- Many drugs act by mimicking or antagonizing the actions of endogenous ligands that regulate flow of ions through excitable membranes via their activation of receptors.
- These receptors are directly coupled (no second messengers) to ion channels.
- For example, the nicotinic receptor for Acetylcholine protein (in nervous system) is coupled to a Na+/K+ ion channel. The receptor is a target for many drugs, including nicotine, choline esters etc.

11

<text><list-item><list-item>

4. Cyclic GMP and Nitric Oxide Signaling

- cGMP is a second messenger in vascular smooth muscle that facilitates dephosphorylation of myosin light chains, preventing their interaction with actin and thus causing vasodilation.
- Nitric oxide (NO) is synthesized in endothelial cells and diffuses into smooth muscle.
- NO activates guanylyl cyclase, thus increasing cGMP in smooth muscle.
- · Vasodilators in synthesis of NO by endothelial cells.



igand-binding

lomaini Iransmembrane

domain

domain

ENZYME

RECEPTOR

Intracellular active enzyme!

5. Receptors That Function as Enzymes or Transporters

1. Receptors That Function as Enzymes or Transporters

• Examples of drug action on transporter systems include the inhibitors of reuptake of several neurotransmitters, including dopamine, GABA, norepinephrine, and serotonin.

2. Receptors That Function as Transmembrane Enzymes

- · These receptors mediate the first steps in signaling by insulin and growth factors.
- Binding of the ligand causes conformational changes (e.g., dimerization) so that the tyrosine kinase domains become activated, ultimately leading to phosphorylation of tissue-specific substrate proteins.

1. 3. Receptors for Cytokines

- · These include the receptors for erythropoietin, somatotropin, and interferons.
- Their receptors are membrane spanning and on activation can activate a distinctive set of cytoplasmic tyrosine kinases Janus kinases [JAKs].
- JAKs phosphorylate signal transducers and activators o f transcription (STAT) molecules.
- STATs dimerize and then dissociate and modulate gene transcription.



Lecture 8 Pharmacokinetics of drugs and medicines

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Absorption Time to peak Cmax Peak level · Concerns the processes of entry of a drug into the systemic Plasma drug concentration circulation from the site of its administration. Minimum effective · The determinants of absorption are those described for concentration drug permeation. Elimination • Intravascular administration (e.g., IV) does not involve absorption, and there is no loss of drug. Time tmax ac Duration of action-Onset of activity • Bioavailability = 100% (discussed in previous lecture) C_{max} = maximal drug level obtained with the dose. t_{max} = time at which C_{max} occurs. Lag time = time from administration to appearance in blood. • With extravascular administration (e.g., per os [PO; oral], intramuscular [IM], subcutaneous [SC], inhalation), less Lag time = time from administration to appearance in biod Onset of activity = time from administration to blood level reaching minimal effective concentration (MEC). Duration of action = time plasma concentration remains greater than MEC. Time to peak = time from administration to C_{max} . than 100% of a dose may reach the systemic circulation because of variations in bioavailability.

Figure I-1-4. Plot of Plasma Concentration Versus Time

5



Distribution

- Drug + Protein \longrightarrow Drug-Protein Complex (Active, free) (Inactive, bound)
- · The processes of distribution of a drug from the systemic circulation to organs and tissue.
- Conditions affecting distribution include:
- Under normal conditions, protein-binding capacity is much larger than is drug concentration. Consequently, the free fraction is generally constant.
- Many drugs bind to plasma proteins, including albumin, with an equilibrium between bound and free molecules (recall that only unbound drugs cross bio membranes).
- Competition between drugs for plasma protein-binding sites may increase the "free fraction;' possibly enhancing the effects of the drug displaced. Example: sulfonamides and bilirubin in a neonate.
- Special Barriers to Distribution:
- Placental-most small molecular weight drugs cross the placental barrier, although fetal blood levels are usually lower than maternal. Example: propylthiouracil (PTU) versus methimazole
- · Blood-brain-permeable only to lipid-soluble drugs or those of very low molecular weight. Example: levodopa versus dopamine
















Problems and answers				
Example:	Pharmacokinetic Problems Solved			
What is the estimated IV LD of <i>Drug A</i> necessary to achieve a concentration of 10 $^{mg}/_{L}$ in a 55 year old, 60-kg male? The V _d of <i>Drug A</i> = 0.5 $^{L}/_{kg}$	 Following the administration of a 250 mg intravenous bolus dose of a drug, the drug- plasma concentration immediately after injection was found to be 17 mcg/mL. Calculate the apparent volume of distribution. 			
C vV	Answer:			
$ID = \frac{O_p \wedge V_d}{D}$	Volume of distribution = $Dose/C_0(conc. at time 0)$			
C _p = desired plasma concentration	= 250/17			
	= 14.7 L			
$LD = \frac{(10 \text{ mg}_{1})* [(0.5 \text{ L}_{kg}) (60 \text{ kg})]}{(60 \text{ kg})}$	Note: 1mcg/ml = 1mg/L			
F Problem 5	2) A patient received a 300 mg dose of an antibiotic by intravenous bolus injection at 6 A.M. At 10 A.M. the concentration of drug in the body was 2.4 mg/mL. If the apparent volume of distribution of this drug is known to be 37 L, calculate the amount of drug in the body at 10			
• For a drug that has an initial plasma concentration of 120 mg/L and a half life of 2 hours, what would the plasma concentration he 12 hours	A.M.			
after the initial concentration?	Answer:			
• A. 15 mg/L	Amount of drug in body = V _d x plasma conc. at that time			
• B. 112.5	= 37 x 2.4			
• C. 7.5 mg/L	= 88.8 mg			
• D 60	5			

Lecture 9 Pharmacokinetics and drug stability

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aA + bB → cC + dD Equilibrium Constant: $K_c = \frac{[C]^c [D]^d}{[A]^a [B]^b}$ (A), (B), (C), and (D) are concentrations of A, B, C, and D respectively

ChemistryLearner.com

Law of Mass Action

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• Rate, order and molecularity

- kinetics is built of the law of mass action.
- This states that the rate of a chemical reaction (i.e. the speed of the reaction or, simply, how fast it is) is proportional to the active masses of the reacting substances.
- Active mass is a complicated term to measure, but, fortunately, if the solutions in question are dilute, the active mass may be replaced by con- centration, which is much easier to handle.
- If the concentration of a solute is greater than about 0.1 mol L⁻¹, significant interactions arise between the solute molecules or ions. In cases like this, effective and measured concentrations are not the same and use must be made of activity instead of concentration.









Rate equations and first-order reactions

- Differential rate equations like the ones above are not much use to the practising chemist, so it is usual to integrate the rate equation to obtain more useful expressions.
- In this reaction, compound A reacts to form products. At the start of the reaction (time 0) the concentration of A is equal to a mol L-1, while the concentration of products will be zero (since the reaction has not started).
- At some later time, t, the concentration of products has increased to x mol L-1 and as a result the concentration of A has fallen to (a x) mol L-1. This can be represented mathematically as
- The rate constant, *k*, is a very important measure of a reaction rate and has the dimension of time-1 for a first-order process

At time = t,

[A] = (a - x) and [products] = x

From the law of mass action, the rate of reaction is proportional to [A]. If we rewrite 'rate' as dx/dt (i.e. the rate of production of x with respect to t), and substituting (a - x) for [A], then

 $\frac{dx}{dt} \propto (a - x)$ and so $\frac{dx}{dt} = k(a - x)$

where \boldsymbol{k} is the constant of proportionality. This expression can be integrated to give

$$\int \frac{\mathrm{d}x}{(a-x)} = \int k \mathrm{d}t = k \int \mathrm{d}t$$
$$-\ln(a-x) + c = kt$$

where ln represents the natural (base e) logarithm. To find c, recall that at t = 0, x = 0; therefore,



Half-life and Shelf-life

- The half-life is defined as the time taken for the concentration of reactant to fall to half its original value:
- For the first order reactions only is independent of concentration

$$\ln \frac{a}{(a-x)} = kt$$
$$\ln \frac{a}{(a-\frac{1}{2}a)} = kt_{\frac{1}{2}}$$
$$\ln 2 = kt_{\frac{1}{2}}$$
$$t_{\frac{1}{2}} = \frac{0.693}{k}$$

- The shelf-life (*t*90) of a pharmaceutical product is the length of time the product may safely be stored on the dispensary shelf before significant decomposition occurs.
- This is important since, at best, drugs may decompose to inactive products; in the worst case the decomposition may yield toxic compounds.
- The shelf-life is often taken to be the time for decomposition of 10% of the active drug to occur, leaving 90% of the activity.
- A similar expression to Eq. (10.3) can be obtained by substituting ln(100/90) in place of ln 2 to give

$$t_{90} = \frac{0.105}{k}$$

 $\frac{1}{(a-x)} - \frac{1}{a} = kt$ $\frac{1}{(a-\frac{1}{2}a)} - \frac{1}{a} = kt_{\frac{1}{2}}$ Therefore, $\frac{1}{\frac{1}{2}a} - \frac{1}{a} = kt_{\frac{1}{2}}$

 $\int \frac{\mathrm{d}x}{(a-x)^2} = \int k \,\mathrm{d}t \qquad \qquad \frac{1}{a} = kt_{\frac{1}{2}}$ e. $t_{\frac{1}{2}} = \frac{1}{ak}$

Hence,

$$\frac{1}{(a-x)} + c = kt$$

At t = 0, x = 0; therefore 1/a + c = 0 and c = -1/a to give

$$\frac{1}{(a-x)}-\frac{1}{a}=kt$$

Second-order reactions $2A \rightarrow \text{products or } A + B \rightarrow \text{products}$

- Equation (10.5) is the equation of a straight line of the type y c = mx; so a plot of 1/(a x) against *t* yields a straight line of slope *k*, with an intercept on the vertical axis of 1/a.
- Equation (10.5) is valid for second-order reactions in which the concentrations of the reactants are equal.
- A general second-order equation may also be derived that will apply to reactions of the type A + B → products when [A] does not equal [B],
- The term *k* is, again, the rate constant for the reaction, but in a second-order process *k* has dimensions of concentration-1 time-1.
- The relationship between the half-life and the secondorder rate constant, *k*, for initial equal concentrations of reactant can be found by substituting t = t 1/2

10

Zero-order reactions

- There are some reactions in which the rate of the reaction is independent of the concentration of the reactants but does depend on some other factor, such as the amount of catalyst present.
- These reactions are termed *zero-order* reactions, and rate equations can be derived as follows:
- In zero-order reactions, the amount of product formed varies with time; consequently, the amount of product formed after 20 minutes will be twice that formed after 10 minutes.
- Reactions that follow zero-order kinetics are quite rare, but they do occur in solid-phase reactions such as release of drug from a pharmaceutical suspension.

$$\frac{\mathrm{d}x}{\mathrm{d}t} = k[\mathrm{A}]^0$$

Therefore,

$$\int \mathrm{d}x = \int k \, \mathrm{d}t$$

which gives

$$x = kt + c$$



Tutorial example

of reactants, the f	ollowin	g data i	vere ob	ained:		
Time (s)	0	89	208	375	625	803
$[Aspirin] (mol L^{-1})$	1.6	1.4	1.2	1.0	0.8	0.7
1 The order of by inspection it r	of a ch nust be	emical determ	reaction	canno	t be de tally. In	termi praci
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where (a - x) is the concentration of each reactant at time *t*, and a plot of 1/(a - x) vs *t* should yield a straight line of slope *k*.

This plot was carried out and a straight line was obtained with a slope of 1.0×10^{-3} . This proves that the reaction is second order with a rate constant, $k=1.0\times 10^{-3}~({\rm mol~L^{-1}})^{-1}~{\rm s}^{-1}$.

 $\frac{1}{(a-x)} - \frac{1}{a} = kt$

Problems and answers

Problems

Q10.1 Determine the first-order rate constant for the hydrolysis of acetyl- β -methylcholine at 85 °C from the information given below.

$[Drug] (mg mL^{-1})$	9.35	7.45	4.52	3.46	1.26	0.90
t (days)	0.08	0.75	1.96	2.96	5.75	6.75

Lecture 10 Medicinal chemistry: the science of rational drug design

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How do drugs work?

- Here, we will introduce the subject of *medicinal* chemistry, which may be defined as the planning and synthesis of new chemical entities for the treatment of disease.
- Medicinal chemistry also includes aspects of molecular pharmacology (i.e. drug action at the molecular level), which, in turn, leads to the holy grail of structureactivity relationships.
- It looks at....How do drugs work?, Where do drugs come from?' and 'Why do we need new drugs?'



How do drugs work?

- Enzyme inhibition
- Enzymes are soluble proteins which function as biological catalysts which increase the rate of chemical reactions without being consumed in the process.
- Most chemical reactions are equilibrium reactions and the presence of the enzyme allows the reaction to reach equilibrium more rapidly than it would in the absence of the enzyme.
- Enzymes do not alter the *position* of the equilibrium, but they do affect the speed with which the reaction reaches equilibrium.
- They do this by forming a high-energy *transition state*, an unstable intermediate formed from the reactants, which decomposes to yield the products of the reaction.





Cyclo- oxygenase forms prostaglandins

- Cyclo- oxygenase, for example, is the enzyme responsible for the formation of inflammatory mediators such as prostaglandins (e.g. prostacyclin) and thromboxane.
- These mediators increase the pain and inflammation associated with minor injury and trauma through vasodilatation, platelet aggregation and release of additional inflammatory mediators.
- Inhibition of the enzyme by drugs such as aspirin, paracetamol or ibuprofen (see Fig) reduce the synthesis of the inflammatory mediators and reduce pain, swelling, inflammation and fever.
- Recent research has shown that there are three isoenzyme variants of cyclooxygenase, known as COX-1, COX-2 and COX-3. Established NSAIDs inhibit all variants of the enzyme.



5

Hypertension inhibitors · Angiotensin-converting enzyme (ACE) is an enzyme produced in the Captopril kidney. **EtOO** • it catalyses the conversion of a short peptide of 10 amino acid residues (a decapeptide), called angiotensin I, into an octapeptide (eight amino acid HOO residues), called angiotensin II. Enalapril • This latter peptide is a very potent vasoconstrictor that rapidly increases blood pressure. · Inhibition of ACE by drugs lowers blood pressure and is used in the treatment of hypertension and congestive heart failure. Lisinopril · Commonly prescribed ACE inhibitors include captopril, enalapril and lisinopril. Structures of captopril, enalapril and lisinopril.

Trimethoprim inhibits DHFR

- Dihydrofolate reductase (DHFR) is an enzyme that catalyses the reduction of dihydrofolic acid to tetrahydrofolic acid, an essential component for the synthesis of purine bases in DNA and certain amino acids.
- Inhibition of DHFR results in a decrease in DNA synthesis and a slowing down of cell division.
- Drugs such as trimethoprim (an antibacterial) achieve their therapeutic action by inhibition of this enzyme.
- Trimethoprim is more active against bacterial DHFR.





The β-adrenoceptor is a G-protein-coupled receptor

- The β -adrenoceptor is a G-protein-coupled receptor found in the cell membranes of many cells.
- Activation of the receptor by adrenaline (epinephrine) results in a response such as an increase in heart rate, an increase in blood pressure etc.
- Blockade of the receptor by drugs like propranolol is useful in the treatment of conditions.
- The propranolol is the natural agonist of the β -adrenoceptor.





Where do drugs come from?

- Has Mother Nature got there before us and done all the hard work herself?
- Examples of a few of the important drugs derived from natural sources,
- local anaesthetics, e.g. lidocaine, derived from cocaine
- narcotic analgesics, e.g. morphine, derived from opium
- antimalarials, e.g. chloroquine, derived from *Cinchona* ACE inhibitors, e.g. captopril, derived from snake venom.
- antibiotics, e.g. penicillin, derived from microorganisms.
- The role of the medicinal chemist is to identify the lead compound and modify it chemically to produce more effective treatments.

Lecture 11 Protein-Ligand docking

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Introduction to proteinligand docking

- Protein-ligand docking is a molecular modelling technique.
- The goal of protein–ligand docking is to predict the position and orientation of a <u>ligand (a small molecule,</u> chemical molecule) when it is bound to a <u>protein</u> receptor or enzyme.
- For example ligand: Paracetamol or any.
- Example of enzyme or protein: lipase, globulin etc.







ΔG_o calculation

- In analogy with any spontaneous process, protein-ligand binding occurs only when the change in Gibbs free energy (ΔG) of the system is negative when the system reaches an equilibrium state at constant pressure and temperature.
- The binding free energy can be calculated using the rate constants k_{on} and $k_{\text{off}}\,as$
- $\Delta G=G$ bound-G unbound =-kTln KeqC₀ = -kTln C₀k_{on}/k_{off}
- where K_{eq} is the binding equilibrium constant, C_0 is the reference concentration of 1 mol/L, k is Boltzmann's constant and T is the temperature in Kelvin

ΔG_o calculation reflect the various contributions to binding.

- The ΔG values on the right of the equation are all constants.
- ΔG_0 is a contribution to the binding energy that does not directly depend on any specific interactions with the protein
- The hydrogen bonding and ionic terms are both dependent on the geometry of the interaction, with large deviations from ideal geometries (ideal distance R, ideal angle α) being penalized.
- The lipophilic term is proportional to the contact surface area (Alipo) between protein and ligand involving non-polar atoms.
- The conformational entropy term is the penalty associated with freezing internal rotations of the ligand. It is largely entropic in nature. Here the value is directly proportional to the number of rotatable bonds in the ligand (NROT).

