# CHEM 191 Chemical Reactions in Aqueous Solution

**Module 1 Lecture 9** 

## Molecules in Biological Environments

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#### **Module 1 Lecture 9**

#### **Learning objectives**

- Recognise the structural features of a molecule which makes it polar or non-polar
- Understand how water solvates ions and polar molecules
- Understand how the pH of the solution determines the ionisation state of a molecule
- Understand how non-polar molecules behave in water
- · Understand the structure and fundamental roles of membranes

#### Polar vs non-polar molecules

A **polar molecule** is one which forms 'favourable' interactions with water molecules. These would include molecules with polar bonds and molecules with charged regions. Polar molecules are also called **hydrophilic** (water loving)

eg  $CH_2OH$   $H_3C$  O  $H_3C$  O  $H_3C$  O

Water is itself a polar molecule – "like dissolves like"

Useful fact to remember alcohols do **NOT** ionise

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#### Polar vs non-polar molecules

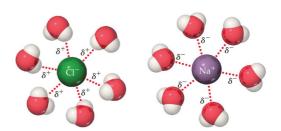
A non-polar molecule is one which does not have any polar bonds or charged parts.

Non-polar molecules will not dissolve in water but will dissolve in other non-polar solvents. Non-polar molecules are also called **hydrophobic** (water hating).

#### Water is a polar molecule

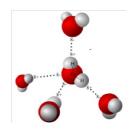
Water - principal component of most cells

Polar nature of water enables electrostatic interaction with other charged molecules - hydration



Water molecules participate in **Hydrogen Bonding** to themselves

Hydrogen Bonding is a very powerful driving force



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#### **Hydrogen Bonding**

Various types of hydrogen bonding are found in biological molecules.

Solvation by water

Within proteins

Within DNA and RNA

#### **Ionisable Functional Groups**

Many molecules (or parts of molecules – **functional groups**) found in biological systems can exist in both charged and uncharged forms.

For example

CH<sub>3</sub>COOH and CH<sub>3</sub>COO-

The ionisation status is influenced by

- 1. The pH of the aqueous environment
- 2. The propensity of particular ionisable functional groups to ionise (fundamental physical property) ie the  $pK_a$

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#### **Ionisable Functional Groups**

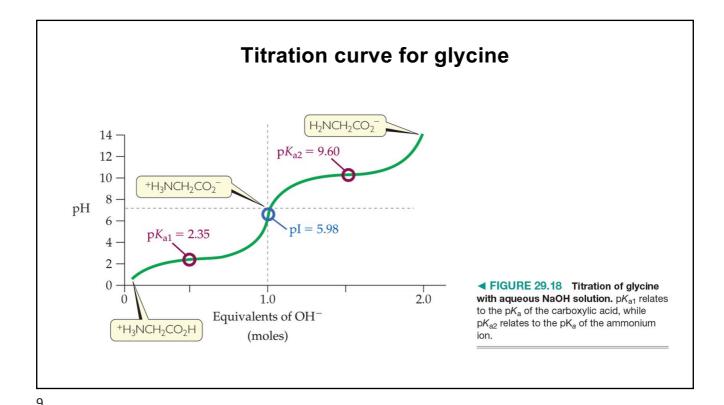
Recall the Henderson-Hasslebach equation for buffers – when the concentrations of the weak acid and the conjugate base are the same,  $pH = pK_a$ 

$$pH = 4.74 + log \frac{(0.100)}{(0.100)} = 4.74$$

Another way to think about this is that

 $pK_a = pH$  of environment in which half (50%) of the functional groups of the molecule are ionised

- >1 pH unit below pK<sub>a</sub> ~90% molecules protonated
- >1 pH unit above pK<sub>a</sub> ~90% molecules deprotonated



#### Ionisable Functional Groups – amino acids

Ionisation possibilities for an amino acid eg alanine

An α-amino group is *protonated* when charged (+ve)
An α-carboxyl acid group is *deprotonated* when charged (-ve)

#### Ionisable Functional Groups – amino acids

Side Chains of some amino acids also contain ionisable functional groups, so their ionisation state depends on the pH of the solution as well.

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#### Ionisable Functional Groups – proteins

In a protein (a long chain of amino acids bonded together), most of the  $\alpha$ -amino and  $\alpha$ -carboxyl groups on amino acids are involved in these bonds and not available for ionisation.

Net charge on such molecules then relies on the ionisable side chain groups (the R groups) and the N- and C-termini (at the end of the chain)

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At physiological pH

#### **Ionisable Functional Groups – buffers**

- To function properly biological systems require molecular components to be in a particular state of ionisation and charge for the evolved purpose
- Variations in pH in the aqueous environment therefore need to be minimised by using buffers
  - The blood buffering system utilises bicarbonate proteins and other molecules also contribute
  - The intracellular buffering system utilises phosphate proteins and other molecules also contribute
- Also, some specialised cell environments may vary more widely in pH

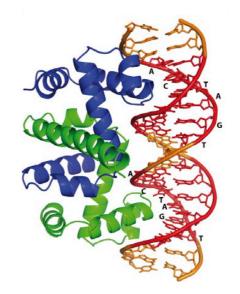
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#### **Ionisable Functional Groups**

Many ionisable groups on macromolecules contribute to charge distribution and influence the shape and nature of interactions of the molecule.

For example,

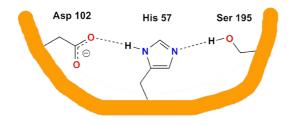
DNA is negatively charged. For a protein to successfully interact with the DNA, the protein must be positively charged.

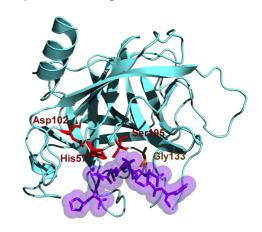


#### Ionisable Functional Groups – enzymes

The ionisation state of amino acid side chains is also crucial for the operation of some enzymes

Eg the active site of Chymotrypsin, a protease (and enzyme which breaks down proteins) contains a **catalytic triad** of amino acid side chains.





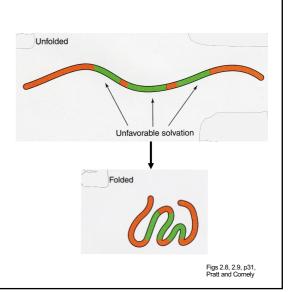
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#### Non-polar molecules in water

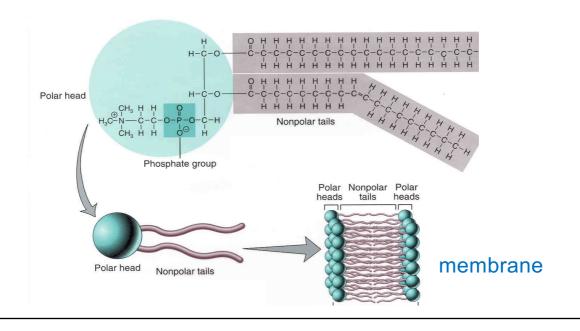
Water molecules will attempt to **hydrate** a nonpolar molecule in an aqueous environment by forming a constrained *hydration shell* around the molecule

Non-polar molecules aggregate together in aqueous solution

Non-polar regions of macromolecules (eg proteins) "hide" away from aqueous environment



#### Molecules with Polar and Non-polar regions - Phospholipids



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#### **Membranes**

Biological membranes provide non-aqueous barriers in aqueous environments in cells

Polar head groups of lipids arranged in bilayers interact with the aqueous environment

The non-polar membrane core formed by the lipid tails effectively constrains molecules to a particular location ie either inside or outside the membrane-bound compartment

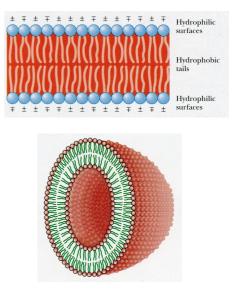


Fig 8.10, p192, Campbell and Farrell 5th ed. Fig 2.12, p33, Pratt and Cornely

#### **Membranes**

- Lipid membranes are heterogeneous
- The lipid composition varies
- The membrane is a dynamic structure
- Other molecules associate with or are inserted in the membrane

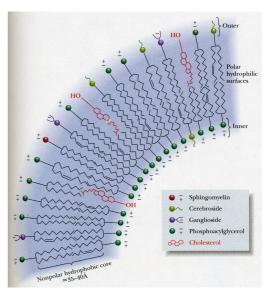


Fig 8.11, p193, Campbell and Farrell 5th ed.

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#### **Membranes**

Other molecules (eg proteins) are inserted in membranes to enable:

- (i) reception and communication of signals
- (ii) transfer of molecules
- across the membrane from one cell compartment to another

The membrane may divide a cell, or an organelle within a cell

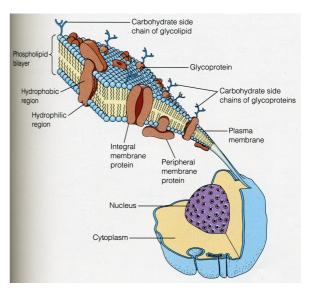


Fig 10.10, p325, Mathews, van Holde and Ahern 3rd ed.

### **Membrane-bound proteins**

A variety of particular amino acid sequence arrangements have evolved for insertion in membranes (eg proteins forming pore structures)

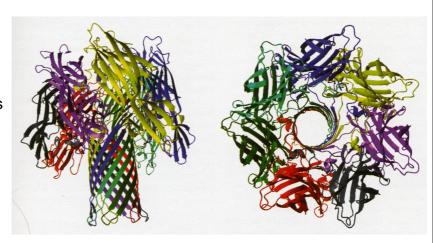


Fig 5.8, p113, Fig 5.31, p131, Whitford

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#### \* Homework \*

#### **No Problems Today**