CHEM 191

Module 4

Structures and reactions of biological molecules

Lecture 7

Peptides – structure and function

Lecturers: Dr Andrea Vernall

Dr Eng Wui Tan

andrea.vernall@otago.ac.nz

ewtan@chemistry.otago.ac.nz

1

Module 4 Lecture 7 Learning objectives

Learning Objectives:

- Plan the synthesis of a target dipeptide from two amino acids, including use of 'protecting groups'
- Recognise that amino acids and peptides are drawn/represented with the N-terminus at the lefthand side of the page
- Understand why amide bonds have restricted rotation and the implications of this on peptide shape
- Recognise that hydrogen bonds and disulfide bridges both influence peptide shape. (+ recap thiol
 oxidation from Lecture 1)

Amide bonds

- In lecture 6 we learnt all about amino acids
- Amino acids join together to make peptides, via amide bonds (called peptide bonds in a peptide)

3

Formation of amide bonds

A carboxylic acid reacts with an amine

Amide bond formation

• This is a type of **nucleophilic acyl substitution** (Lecture 5) but a 'coupling reagent' is required.

..... Why do we need a coupling reagent?

Δ

Formation of amide bonds

- Why do we need a peptide coupling reagent to make an amide?
 - o 1. the carboxylic acid is not a great electrophile (Lecture 5), and
 - o 2. if 'just' an amine and carboxylic acid are combined this happens -

$$OH + H_2N$$
 Acid-base reaction

Why won't adding an acid (H⁺) help like we saw with ester formation?

First step of ester formation

Because with an amine, this will happen

$$H_2N \longrightarrow H_3N \stackrel{\oplus}{\smile}$$

And there is no longer an amine nucleophile

5

Formation of amide bonds

 We are not covering the details of what 'coupling reagents' are used for peptide synthesis in CHEM 191, just recognizing that a 'coupling reagent' is required

$$OH + H_2N$$
 coupling H H_2O H H_2O

 The coupling reagents essentially do similar chemistry that we have already learnt in Lecture 5 – if we turn the carboxylic acid into something more reactive, e.g.

OH O
$$+$$
 H_2N $+$ H_2O acid chloride easily reacts with amines

Synthesis of peptides

 Since an amino acid contains an amine and a carboxylic acid, if two unprotected amino acids are reacted, then four dipeptide products are possible

- · Peptides are always written with the N-terminus at the LHS of the page
 - A-G is a different compound to G-A

7

Synthesis of peptides

• Solution to making only 1 target product – protect the functional groups that you don't want to react. (i.e. temporarily block the reactivity in that position).

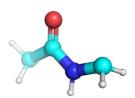
E.g. if the target peptide is ala-gly

Peptide bond structure

• The lone pair of electrons of the amide nitrogen is in resonance / the nitrogen lone pair of electrons is delocalised towards the carbonyl group oxygen

- This means that peptide bonds in peptides are

 - 2. Rigid and planar, due to the delocalised electrons and restricted rotation of the O-C-N bonds.



q

Peptides

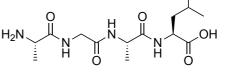
- · Peptides are polymers of amino acids
- · Can be named according to the number of monomers in the peptide -

2 amino acids
 3 amino acids
 many amino acids
 polypeptide

Is it a peptide or protein??

The word 'protein' is generally used to describe peptides than are greater than 50 amino acids long

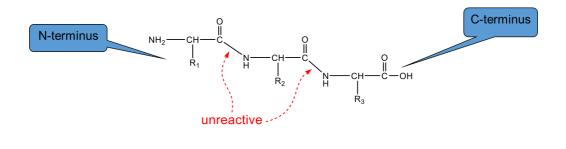
- Peptides have a directional sense, like amino acids do
 - As for amino acids, always write peptides with the N terminus on the left



Ala-Gly-Ala-Leu

Peptide properties

- Peptide properties are governed by the sequence of amino acid units, this is called the **primary structure**
 - o Different amino acids have different **side chains**, the properties of these **side chain** groups (e.g. ionisation at certain pHs, hydrogen-bonding) influence peptide properties.

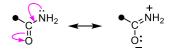


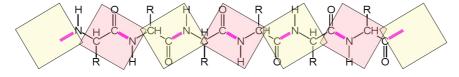
11

Peptide properties

• Because of the restricted rotation of the O-C-N bonds







Bond rotation is not easily possible about the amide bonds

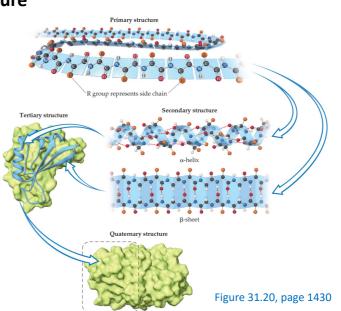
Sequential α -carbons are usually in a *trans* relationship

Peptide, protein shape and structure

tertiary structure - how secondary structural elements fit together

quaternary structure - how proteins or independent peptide chains come together





13

Why to peptides adopt secondary structures?

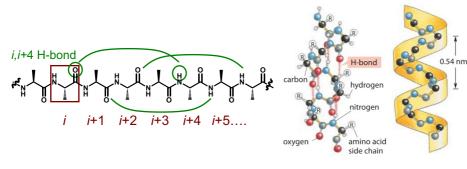
Page 1431

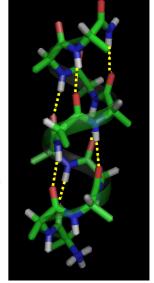
 In an aqueous environment, the chain will adopt a conformation to expose the polar side chains (hydrophilic groups) and bury the non polar side chains (hydrophobic groups) i.e. maximise favourable non-covalent interactions

 Hydrogen bonding between different amide bonds also stabilises secondary structural structures

Secondary structure example – alpha helix

- The NH from an amide hydrogens bonds with the CO of a different amide 4 amino acids along the chain
- This same pattern of hydrogen bonds repeated along the peptide gives the stabilised helical secondary structure

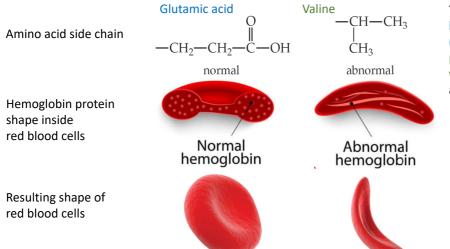




15

Even one amino acid change can change the shape of a protein

• Sickle-cell anaemia is a genetic disease where one glutamic acid in the protein haemoglobin is replaced with the amino acid valine.



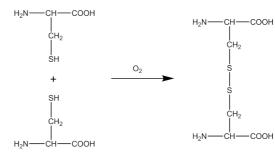
The change from the hydrophilic glutamic acid (CO₂-at pH 7.4) to the non-polar, more hydrophobic valine causes the protein to aggregate

Page 1420

Disulfide bridges

• Depending on the protein, disulfide bond (or bridges) can be defined as stabilising both secondary or tertiary peptide/protein secondary structures.

- The amino acid cysteine contains a thiol functional group on it's side chain.
- Even under just mild air oxidation conditions, two cysteines can react in an oxidation reaction to give a disulfide bond



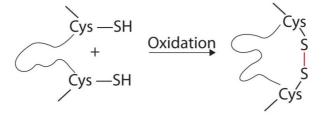
Two cysteines, each with a thiol

Disulfide bond

17

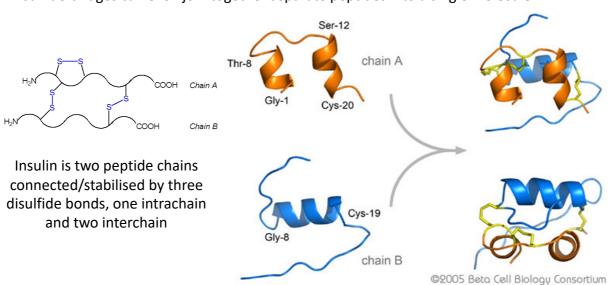
Disulfide bridges

- A peptide may contain the amino acid cysteine in several places.
- If a disulfide bridges forms, this can even link/bring together otherwise remote ends of the peptide.



Disulfide bridges – insulin example

• Disulfide bridges can even join together separate peptides into a single molecule



19

Homework *

31.21, 31.23, 31.25, 31.62, 31.63(a), 31.64, 31.67(c, harder)(d)