Alanine (*ala*) and lysine (*lys*) are two amino acids with the structures given below as Fischer projections. The pK_a values of the conjugate acid forms of the different functional groups are indicated.

 $pK_{a} = 2.35$ $pK_{a} = 2.35$ $pK_{a} = 2.18$ $pK_{a} = 9.87$ $H_{2}N - H_{1}$ CH_{3} ala $PK_{a} = 8.95$ $H_{2}N - H_{1}$ $CH_{2}N + H_{2}$ $PK_{a} = 10.53$ lys

Draw the structure of the dipeptide *ala-lys* in its zwitterionic form.



Note that the amine group on the side chain is more basic so it is the one that is protonated.

Would you expect the dipeptide to be acidic, neutral or basic? Give a brief reason for your choice.

Basic. The side chain in lysine is basic whilst that in alanine is neutral.

Estimate the isoelectric point of the dipeptide.

The isoelectric point occurs when the peptide has no overall charge. The dipeptide has 2 amine and one carboxylic acid group left after the formation of the amide bonds.

The zwitterionic form is the third form drawn overleaf, occurring between pH = 9.87 and 10.35.

pI is half way between these values: $pI = \frac{1}{2} (9.87 + 10.35) = 10.20$.

ANSWER CONTINUES ON THE NEXT PAGE

CHEM1611





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- Marks
- Alanine $(R = CH_3)$ and lysine $(R = CH_2CH_2CH_2CH_2NH_2)$ are two common amino 6 acids. Using *ala* and *lys* to represent the two amino acids, represent all constitutional isomers of the tripeptide formed from one *ala* and two *lys* units. ala-lys-lys lys-ala-lys lys-lys-ala Comment, giving your reason(s), on whether the tripeptide(s) will be acidic, neutral or basic in character. The alanine side chain is neutral whilst the lysine side chain is basic. The tripeptide will therefore be basic. The pK_a values of lysine are 1.82 (α -COOH), 8.95 (α -NH₃^{\circ}) and 10.53 (side chain). What is the value of the isoelectric point of lysine? Fully protonated, lysine has a –COOH group and two NH_3^+ groups. It has a +2 charge. To get a neutral form, both $=NH_3^+$ groups must be deprotonated. The isoelectric point is thus the mean of the pK_a values for these two groups: $pI = \frac{1}{2}(8.95 + 10.53) = 9.74$ pI = **9.74** Draw the Fischer projection of the zwitterionic form of lysine. $\begin{array}{c} & & & \\ & & & \\ H_2N - & H \\ & & & \\ & & \\ & & \\ & & \\ CH_2CH_2CH_2CH_2NH_3^{\oplus} \end{array}$



THIS QUESTION CONTINUES ON THE NEXT PAGE.



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Marks • Neurontin[®] is a pharmaceutical now widely used for the treatment of nerve pain. The 4 structure of the active ingredient in Neurontin, gabapentin, is shown below. The pK_a value for the carboxyl group is 3.68, whilst the pK_b value for the amine group is 3.30. gabapentin OH NH_2 Explain whether gabapentin can reasonably be described as an amino acid. Gabapentin has both an amine and a carboxylic acid functional group, so it is an amino acid. It is not an α -amino acid (like those found in proteins) as the amino group is not attached to the carbon next to the COOH group. Orally-delivered pharmaceutical agents that contain amine functional groups are often prepared as hydrochloride salts, rather than as free amines. Suggest a reason why gabapentin is not delivered as a hydrochloride salt, illustrating your answer with a suitable diagram. Salt formulations are mainly used to prevent oxidation of the free amine group. The amine group is converted to a quaternary ammonium salt which is more stable. However, gabapentin already exists in a zwitterionic form at normal pH, with the amine group protonated to form the more stable quaternary ammonium ion. Ò OH NH_2 NH₃ ANSWER CONTINUES ON THE NEXT PAGE

Gabapentin was originally synthesised as it was anticipated that it would bind to the same receptors as the neurotransmitter GABA (4-aminobutanoic acid). Draw the structure of GABA. Suggest a reason why it might have been anticipated that gabapentin would interact with GABA receptors, and what form such interactions might take.



GABA (4-aminobutanoic acid).

Both GABA and gabapentin have the same basic features - a four carbon chain with a terminal NH_2 and a terminal COOH group. These functional groups are likely to be involved in receptor binding through interactions such as H-bonding.

ANSWER CONTINUES ON THE NEXT PAGE





Define the terms *primary structure, secondary structure* and *tertiary structure* in relation to proteins. Illustrate your answer with appropriate diagram(s) and by making reference to the representations shown above.

ANSWER CONTINUES ON THE NEXT PAGE.

Marks 4

The peptide links in a protein chain are said to be *resonance stabilised*. Use a diagram to explain what is meant by this term, and indicate one important consequence relating to protein structure and one important consequence relating to the chemistry of proteins.

Resonance occurs when two or more Lewis structures can be drawn for the same compound. In such cases, the true structure is none of those drawn, but rather a weighted average of all of them.



The amide functional group has two major resonance contributors as shown. As a consequence of resonance, the peptide bond is rigid, planar and strong.

Consequence for structure: This rigidity and the charge on the oxygen are ideal for the formation of α -helices and β -pleated sheets via H-bonding.

Consequence for chemistry: The involvement of the N lone pair in resonance, means that the N is unavailable for protonation and is non-basic. The peptide bond is therefore relatively inert.

Modern medicine now uses insulin analogues (where one or more of the amino acid residues has been changed) in the treatment of diabetes. In one such analogue, glargine insulin, the changes have increased the isoelectric point of the enzyme from 5.4 to 6.7, thereby reducing its solubility at physiological pH. Explain how changes in the primary amino acid sequence can alter the pI and solubility of the analogue without altering its interaction with blood-glucose.

Changing surface amino acids (for example by changing charged groups to uncharged or polar groups to non-polar) alters the pI and hence the solubility of the protein.

As long as the residues changed are not near the active site and do not change its shape, the mode of action of the enzyme is not affected.



Marks • The structure of the naturally occurring tetrapeptide His-Phe-Ala-Glu, A, is shown 10 below as the zwitterion. $\begin{array}{c} H_{3}\overset{\oplus}{N}-CH-CO-NH-CH-CO-NH-CH-CO-NH-CH-CO_{2} \\ \downarrow \\ CH_{2} \\ \downarrow \\ CH_{2} \\ CH_{2} \\ CH_{3} \end{array}$ A ĊOOH ١H Give the product(s) obtained when **A** is treated with cold 1 M NaOH solution. —сн—Ё-—Ņ-—n—сн—ё-| | NH₂-CH-Ö -СН- CO_2 ĊH₂ Ĥ Ĥ Ĥ CH₂ CH₃ Θ CO_2 NH Ν Give the Fischer projections of the four L-amino acids in their correct ionic states obtained from the vigorous basic hydrolysis (6 M KOH) of A. co_2^{Θ} $\operatorname{CO}_2^{\Theta}$ $\operatorname{CO}_2^{\Theta}$ CO_2^{\ominus} $\begin{array}{cccc} -H & H_2 N - \stackrel{l}{C} - H & H_2 N - \stackrel{l}{C} - H \\ H_2 & C H_3 \end{array}$ H_2N H_2N -H ĊH₂ CH₂ .NH N



Given that the pK_a of the carboxylic acid group of leucine is 2.32 and the pK_b of the amine group is 4.24, do you expect the classical or the zwitterionic form to predominate when leucine is dissolved in water? In other words, does the following equilibrium lie to the right or left? Show your reasoning.

$$H_2N-CH(CH_2CH(CH_3)_2)-COOH \iff H_3N-CH(CH_2CH(CH_3)_2)-CO_2^{\ominus}$$

The equilibrium for the K_a of the acid group is:

$$H_2N$$
-CHR-COOH \iff H_2N -CHR-COO⁻ + H^+

for which:

$$K_{a(COOH)} = \frac{[H^+][H_2NCHR - COO^-]}{[H_2NCHR - COOH]} = 10^{-2.32}$$

The equilibrium for protonation of the amine group is:

$$H_2N$$
-CHR-COO⁻ + $H^+ \iff H_3N^+$ -CHR-COO⁻

for which:

$$K = \frac{[\mathrm{H}_{3}\mathrm{N}^{+} - \mathrm{CHR} - \mathrm{COO}^{-}]}{[\mathrm{H}_{2}\mathrm{NCHR} - \mathrm{COO}^{-}][\mathrm{H}^{+}]} = \frac{1}{K_{\mathrm{a(NH}_{3}^{+})}} = \frac{1}{10^{-9.76}} = 10^{+9.76}$$

in which $pK_a + pK_b = 14$ has been used.

The equilibrium for formation of the zwitterionic form in the question is:

$$H_2N$$
-CHR-COOH \iff H_2N^+ -CHR-COO⁻ + H^+

for which:

$$K' = \frac{[\text{H}_{3}\text{N}^{+} - \text{CHR} - \text{COO}^{-}][\text{H}^{+}]}{[\text{H}_{2}\text{NCHR} - \text{COOH}]} = \frac{K_{a(\text{COOH})}}{K_{a(\text{NH}_{3}^{+})}} = 10^{-2.32} \times 10^{+9.76} = 10^{7.44} >> 1$$

As the equilibrium constant >> 1, the equilibrium lies far to the right and so the zwitterionic form dominates.

Marks • The neurohormone Tyr-Gly-Gly-Phe-Met (T) known as methionine enkephalin is a 7 naturally occurring peptide which controls pain perception in vertebrates. $-C-NH-CH_2-C-NH-CH-CH$ $CH - C - NH - CH_2 -$ $_{H_3N}^{\oplus}$ — -NH- $CH - CO_2$ ĊH₂ ĊH₂ ĊH₂ Ļ CH2 **(T)** CH₃ ÓН Name the functional groups in (**T**). tertiary ammonium ion, phenol, amide, arene, carboxylate ion, thioether Four amino acids (tyrosine, glycine, phenylalanine and methionine) are obtained on complete acid hydrolysis of (T). Draw the stereoformulas of L-tyrosine and L-methionine in the boxes below. Indicate their absolute configurations using the (R)- and (S)- convention. L-methionine L-tyrosine HO₂C.^b HO₂C, **b** '''''H⊕ ''''' H⊕ 'NH₃ NH₃ с с a я OH Absolute configuration S S Absolute configuration Give the constitutional formula for the product obtained when tyrosine, the

N-terminal amino acid in peptide (**T**), is dissolved in 1 M NaOH solution.







Give the constitutional formulas for the following dipeptides in their zwitterionic states.







The heterocycle present in the sidechain of histidine is imidazole, whose structure is shown on the right. Give the structure of a tautomer of imidazole and state, giving reasons, whether your tautomer is aromatic.



Imidazole is aromatic: there are 6π electrons, the ring is planar and each ring atom is sp² hybridized. The C=C and C=N bonds both contributes 2π electrons. The N-H nitrogen atom has two electrons in a p-orbital which also contribute to the π system.

The lone pair on the second nitrogen are housed in a sp² hybrid direct away from the molecule and do not contribute to the π electrons.

What is the major species present when histidine is dissolved in water at pH 12. The pK_a values of histidine are 1.82 (-COOH), 9.17 (-NH₃^{\oplus}) and 6.04 (sidechain).

