### **CONCEPT: SECONDARY PROTEIN STRUCTURE**

• The type of structure that results from \_\_\_\_\_ of the atoms in the backbone of a protein.

□ Involves the connection between the \_\_\_\_\_ of one peptide with the \_\_\_\_\_ of another.

**EXAMPLE:** Determine which of the following amino acid pairs could potentially perform hydrogen bonding between their respective R groups?

a) Gly and Ser

b) Asp and Glu

c) Val and Leu

d) Asp and Arg

**PRACTICE:** How many hydrogen bonding pairs are possible when the following two peptides interact?

a) 3

b) 5

c) 4

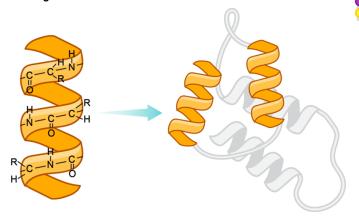
d) 6

e) 1

# **CONCEPT: SECONDARY PROTEIN STRUCTURE**

## Alpha-Helix

- Secondary structures give rise to 2 types of repeating patterns.
- The backbone of a single protein chain into a spiral-like staircase.
  - □ Stabilized by -bonding between *distant* amino acids on the same chain.



Primary

Structure

Secondary

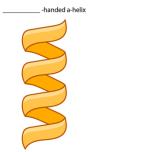
Structure

**EXAMPLE:** Determine which of the following statements represents a 2° structure for a protein.

- a) Creation of peptide bonds.
- b) The attractive force between the H atom of a peptide bond and the O atom of a peptide bond.
- c) Amide bond formation in the creation of an amino acid chain.
- d) Ionic bond formation between the R side chains of alanine and valine.

#### Alpha Helix Spiral Shape

- The spiral-like staircase adopts a right-handed (\_\_\_\_\_\_) shape.
  - □ The hydrogen bonds lie \_\_\_\_\_ the helix and the amino acid R groups lie \_\_\_\_\_ the helix because of spacing.



- The hydrogen bonding of the amide hydrogen with the carbonyl oxygen happens \_\_\_\_\_ residues further on the helix.
  - ☐ The result is every \_\_\_\_ turn of the helix contains on average \_\_\_\_residues

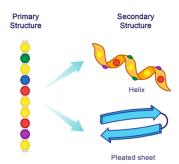
**EXAMPLE:** What is the maximum number of turns for an alpha helix that contains 72 residues?

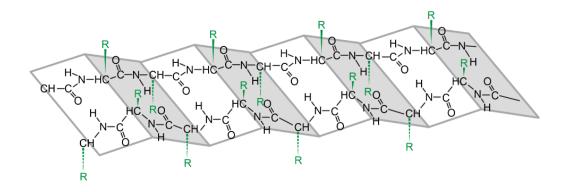
- a) 15 turns
- b) 18 turns
- c) 72 turns
- d) 20 turns
- e) 6 turns

# **CONCEPT: SECONDARY PROTEIN STRUCTURE**

## **Beta-Pleated Sheet**

- Secondary structure consisting of \_\_\_\_\_ or more  $\beta$ -strands oriented side by side.
  - □ Named "pleated" because of their \_\_\_\_\_- structure.
  - $\ \square$  R side chains extend \_\_\_\_\_ or \_\_\_\_ to the  $\beta$ -sheet.





**EXAMPLE:** Which of the following statements is true of  $\beta$ -sheets?

- a) Interchanging between an  $\alpha$ -helix and a  $\beta$ -sheet is a key feature of a primary structure.
- b) Their interior is characterized by hydrogen bonding between amide hydrogens and carbonyl oxygens.
- c) Their interior is characterized by R side chains interactions.
- d) The R side chains extend inward to ensure greater packing of the peptides.

**PRACTICE:** Which of the following statements is true in regard to the peptide strand shown?

- a) The  $\beta$ -sheet defines the primary structure of the peptide strand.
- b) The C-Terminal end possesses an  $\alpha$ -helix.
- c) Along with its  $\alpha$ -helix counterpart, the  $\beta$ -sheet is mainly stabilized by backbone hydrogen bonds.
- d) The  $\alpha$ -helix and the  $\beta$ -sheet are connected together through an ionic bond.

