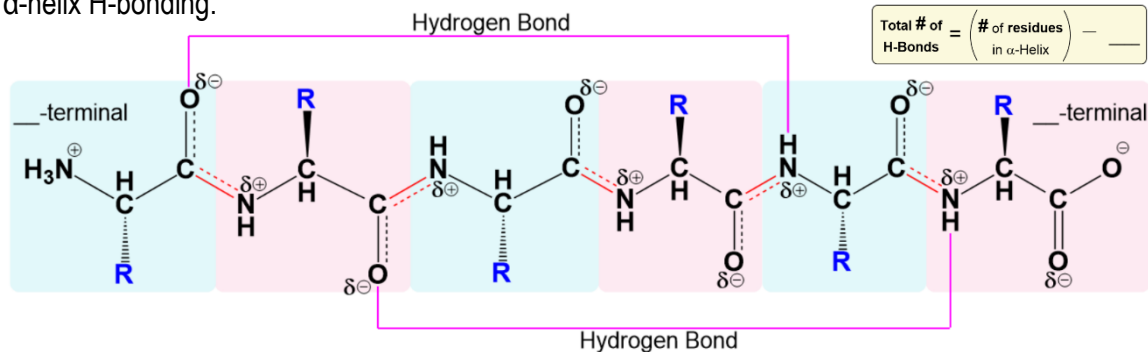


## CONCEPT: ALPHA HELIX HYDROGEN BONDING

- $\alpha$ -helices are stabilized by intrachain hydrogen bonds between N-H and C=O groups in the peptide \_\_\_\_\_.
  - R-group hydrogen bonding is \_\_\_\_\_ involved in  $\alpha$ -helix stabilization.
- Each C=O group of a residue hydrogen bonds with the N-H group \_\_\_\_\_ residues away (residue "X" bonds with X+4).
  - Therefore, first & last *four* residues of an  $\alpha$ -helix do \_\_\_\_\_ fully participate in  $\alpha$ -helix hydrogen bonding.

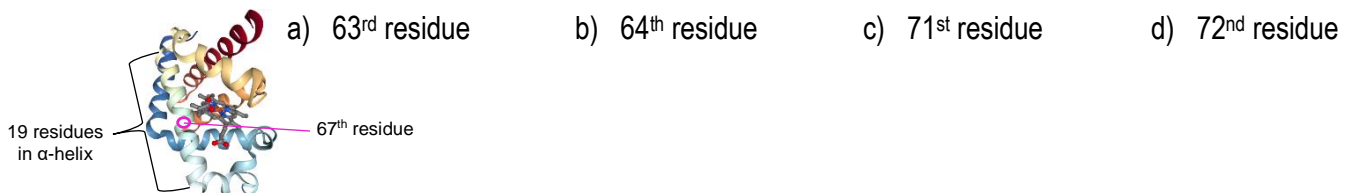
EXAMPLE:  $\alpha$ -helix H-bonding.



EXAMPLE: Which residue does the carbonyl group of the 21<sup>st</sup> residue of a 30 residue  $\alpha$ -helix hydrogen bond to?

- a) 25<sup>th</sup> residue.      b) 26<sup>th</sup> residue.      c) 32<sup>nd</sup> residue.      d) 33<sup>rd</sup> residue.

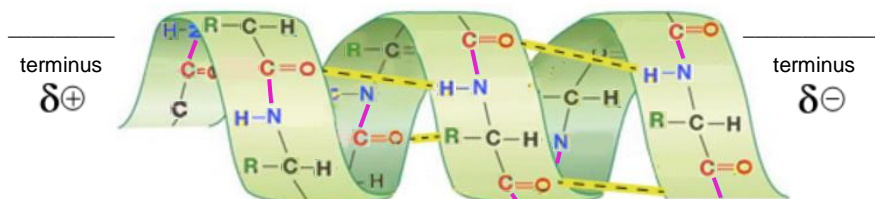
PRACTICE: The oxygen-storage protein myoglobin has 8  $\alpha$ -helices in its single polypeptide chain. Its 67<sup>th</sup> residue is near the center of a 19 residue  $\alpha$ -helix. Which residue does the amino group of myoglobin's 67<sup>th</sup> residue hydrogen bond to?



## $\alpha$ -Helix Net Dipole

- $\alpha$ -helices have an overall net \_\_\_\_\_ due to the direction of polar peptide bonds & intrachain hydrogen bonds.
  - Net electron density is shifted towards the \_\_\_\_\_-terminus.
  - N-terminus of the  $\alpha$ -helix has a net \_\_\_\_\_ charge while the C-terminus has a net \_\_\_\_\_ charge.

EXAMPLE:  $\alpha$ -helix Net Dipole.



PRACTICE: True or False: Negatively charged residues near the  $\alpha$ -helix N-terminal are stabilizing due to its net dipole.

- a) True.      b) False.

**CONCEPT: ALPHA HELIX HYDROGEN BONDING**

**PRACTICE:** Triose phosphate isomerase (TPI) is a crucial enzyme involved in the glycolysis pathway and contains 14  $\alpha$ -helices. Considering the net dipole of the  $\alpha$ -helix, which of the following would be most destabilizing to TPI's structure?

- a) An electric dipole spanning several peptide bonds throughout its  $\alpha$ -helices.
- b) The presence of Glu residues near the N-terminus of its  $\alpha$ -helices.
- c) The presence of Arg residues near the C-terminus of its  $\alpha$ -helices.
- d) The presence of Lys residues near the N-terminus of its  $\alpha$ -helices.

**PRACTICE:** The hemagglutinin protein in influenza virus contains a remarkably long  $\alpha$ -helix with 53 residues.

- A) How long is the  $\alpha$ -helix? \_\_\_\_\_
- B) How many turns does this  $\alpha$ -helix have? \_\_\_\_\_
- C) How many hydrogen bonds are present in this  $\alpha$ -helix? \_\_\_\_\_

**PRACTICE:** Which of the following statements about  $\alpha$ -helices is false?

- a) Myoglobin & hemoglobin  $\alpha$ -helices are right-handed  $\alpha$ -helices.
- b) Each residue of an  $\alpha$ -helix creates a  $100^\circ$  turn of the  $\alpha$ -helix backbone.
- c) The core of an  $\alpha$ -helix is tightly packed with backbone atoms.
- d)  $\alpha$ -helices have an overall macrodipole with a partially positive C-terminus & partially negative N-terminus.
- e) Hydrogen bonds that hold the  $\alpha$ -helix together are about parallel to the axis of the helix.