## **CONCEPT: PROTEIN MOTIFS AND DOMAINS**

<ul> <li>Tertiary protein structure in</li> </ul>	cludes the distribution	of $\alpha$ -helices, $\beta$ strar	nds & turns/loops	s to make	&
Primary Structure (Amino acid+ sequence)	Secondary S (Exhelix,shee		upersecondary (	Structure  _) ertiary Structu	(Independent folding units)
Protein Motifs					
<u>Motifs</u> (or supersecondary structures): specific patterns &			of α-helices, β strands, & turns/loops.		
☐ Have different functions in proteins including providing					
•Several of l	•				
1. Helix-loop-helix		3. βαβ			
<b>EXAMPLE:</b> Motifs.	2				
1	2.	9999	3.	4.	
PRACTICE: Which of the fo	llowing colored region	s in the images belo	w is not an exan	nple of a supers	econdary structure?
a) Greek key.	b) β-meander.	c) βαβ unit.	d) Type	e I β-turn	e) Helix-loop-helix
Protein Domains					
•: combinations of Domains are an e	of motifs that <i>independ</i> xtension to a peptide's	•	•		
●Proteins are often	according to the	structures & function	nal characteristic	cs of their doma	ins.
<b>EXAMPLE</b> : Circle the 3 mos	st obvious domains in	the figure.			

**PRACTICE:** Which of the following statements concerning a Cas9 endonuclease's protein domains is true?

- a) They are a form of its secondary structure.
- c) All its domains consist of separate polypeptide subunits.
- b) They have only been found in eukaryotic proteins.
- d) They retain their shape when separated from the protein.

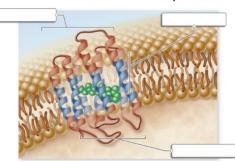
## **CONCEPT: PROTEIN MOTIFS AND DOMAINS**

**PRACTICE:** Which of the following is true concerning the motifs and domains of proteins?

- a) Many domains make up a motif.
- b) Every polypeptide chain is limited to one domain.
- c) Separate proteins with similar domains are likely to have a similar function.
- d) All domains of a protein have the same function.

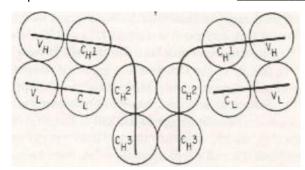
**PRACTICE:** Appropriately label the domains of the membrane embedded protein in the figure below.

- a) Transmembrane domain.
- b) Cytoplasmic domain.
- c) Extracellular domain.



**PRACTICE:** The structure of an immunoglobulin G (antibody) molecule is shown schematically below. The black solid lines depict individual polypeptides and so there are four polypeptides in the quaternary structure of this molecule. Each of the spheres represents a stretch of about 100 amino acids folded independently of the rest of the polypeptide and performs a specific function in the molecule. Therefore, each sphere was given its own individual name (VH, CH1, CH2, etc.). Without knowing any additional details, you can predict that there must be TWELVE \_\_\_\_\_\_\_ in this molecule.

- a) α-helices.
- b) Domains.
- c) Subunits.
- d) Motifs.



PRACTICE: What is the main difference between an endonuclease's domains and its subunits?

- a) Its domains are composed mostly of the  $\alpha$ -helix, while subunits contain both  $\alpha$ -helices and  $\beta$ -sheets.
- b) Its subunits are separate polypeptide chains, while its domains constitute a part of a polypeptide chain.
- c) Its domains do not have secondary structure, whereas its subunits have do.
- d) Its domains are stabilized by hydrogen bonds, while its subunits are stabilized by disulfide bonds.