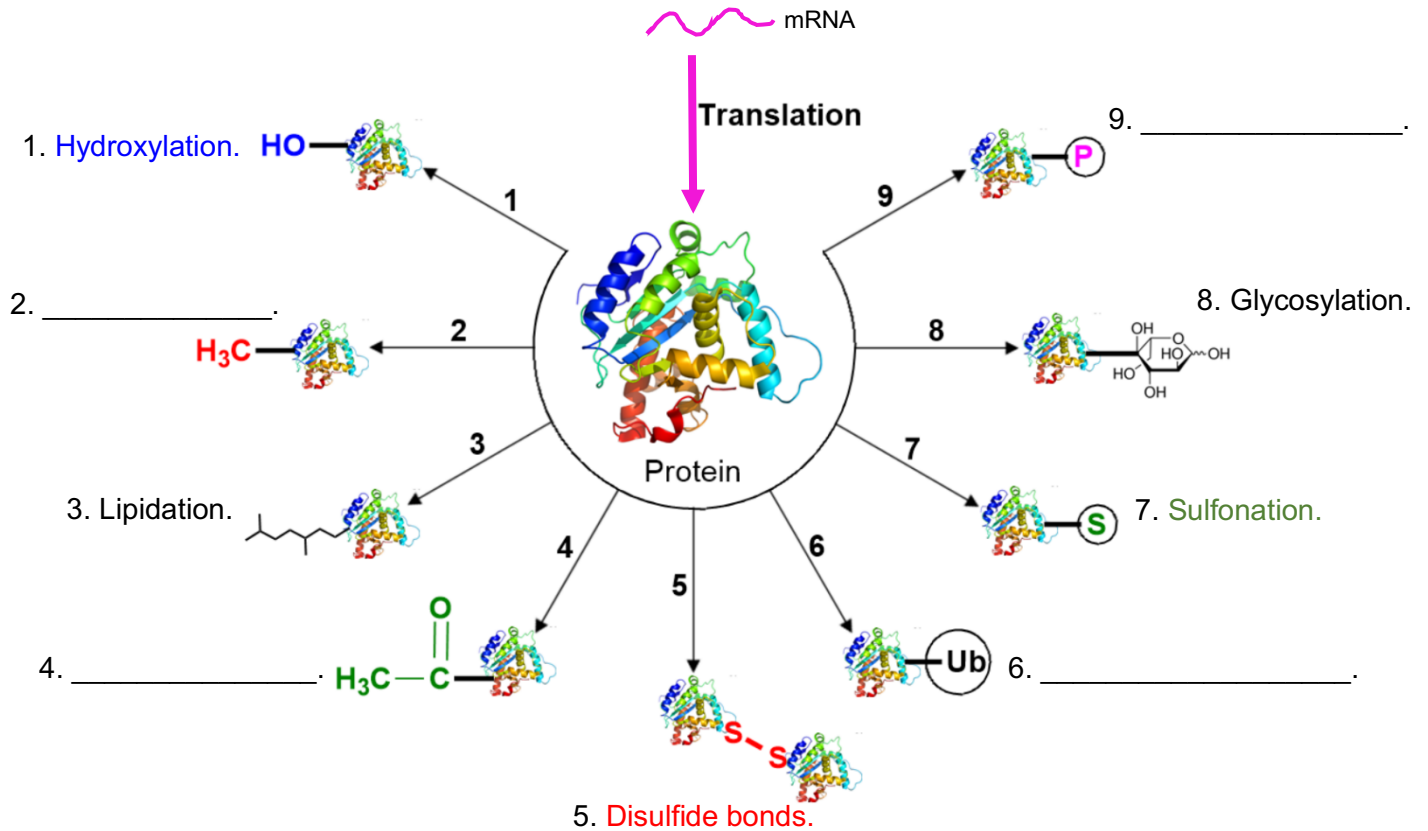


CONCEPT: POST TRANSLATIONAL MODIFICATION

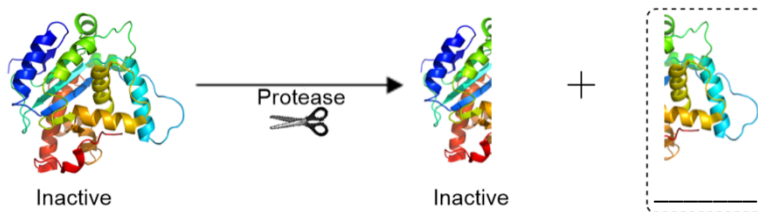
- Recall: *translation* is the cellular process of building _____ using the encoded messages of _____.
- _____-Translational Modifications: protein _____ alterations that control protein activity _____ translation.
- There are *many* types of *post-translational modifications* (PTM), but some of the more common types include:
 - Methylation
 - Acetylation
 - Ubiquitination
 - Phosphorylation

EXAMPLE: Post-Translational Modifications.



Proteolytic Cleavage

- Another type of *post-translational-modification* is _____ cleavage.
 - Proteolytic cleavage*: process of *breaking* _____ bonds between amino acid residues of proteins.
 - Requires proteases or _____, which are *enzymes* that cleave peptide bonds.



PRACTICE: Which of the following would not be a reversible post-translational modification?

- a) Phosphorylation.
- b) Preoteolytic processing.
- c) Alkylation.
- d) Methylation.

CONCEPT: POST TRANSLATIONAL MODIFICATION

PRACTICE: An enzyme's active site has high affinity for a polar charged substrate. How will methylation of the active site affect the reaction rate of the enzyme?

- a) Affinity for the charged substrate will increase thereby increasing reaction rate.
- b) Affinity will remain unchanged, and therefore will be independent of the modification.
- c) Affinity for the substrate will decrease, lowering the reaction rate.

PRACTICE: Histones are proteins that regulate gene expression by binding to DNA and controlling which regions are exposed to be expressed. Histones bind to DNA at a neutral pH via their positively charged lysine residues, but acetylation of histones leads to the dissociation of the DNA-Histone complex. Covalent modification of lysine's ammonium group by acetylation results in an overall _____ charge of the histone binding region.

- a) Positive.
- b) Neutral.
- c) Negative.