

CONCEPT: INDIRECT PROTEIN SEQUENCING VIA GENOMIC ANALYSES

- _____ protein sequencing (via Tandem MS or Edman degradation), is used on *already* extracted/isolated proteins.
 - *Directly* identifies sequence of _____ proteins in a sample; HOWEVER....
- Most protein sequencing data is derived *indirectly* from _____ analyses (translating nucleotide sequences of genes).

Why Obtain Protein Sequences via Genomic Analyses?

- It saves a lot of time! Working with DNA is *easier* than working with proteins.
 - DNA sequencing is significantly _____, cheaper & *more* efficient/informative than direct protein sequencing.
 - Allows us to collect more protein sequencing data faster!

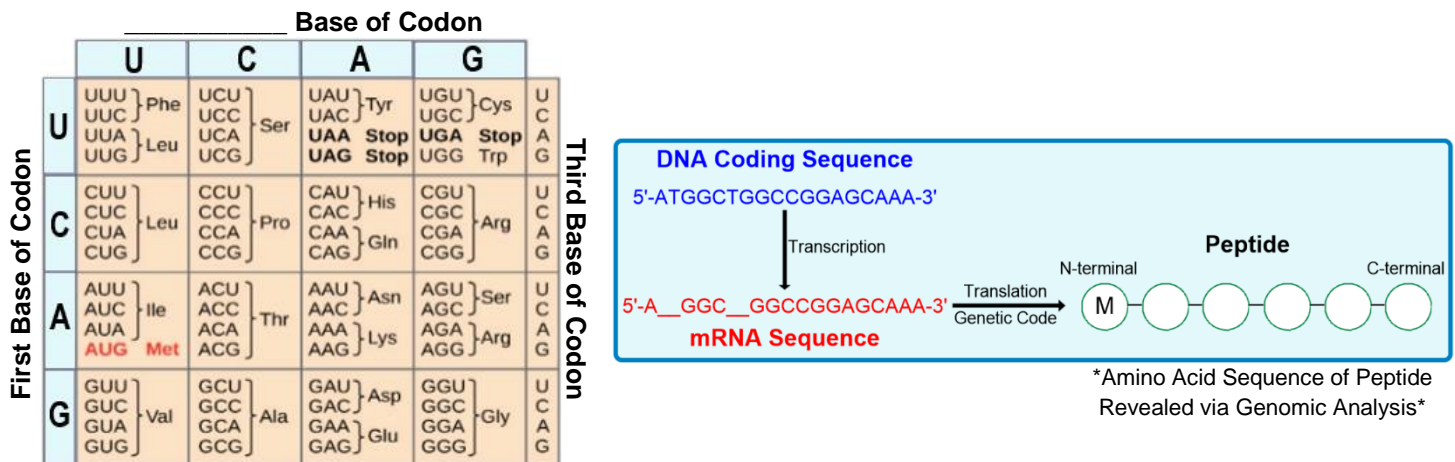
So Why Do We Even Need Direct Protein Sequencing?

- We can't just scrap direct protein sequencing because it has its own advantages!
 - Genomic analyses cannot identify an unknown protein sample on its own.
 - Unlike genomic analyses, Tandem MS can reveal chemically _____ residues (ex. lipoproteins, etc.).

Genetic Code

- Recall: _____ code reveals the connection between codons of nucleic acids & amino acids of proteins.

EXAMPLE: Use the Genetic Code to reveal the peptide sequence.




PRACTICE: Use the genetic code above & the coding DNA sequence below to determine the protein sequence.

5'-ATGGCCTGCGTTCTCAAG-3' →

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PRACTICE: Suppose the sequence below is a template DNA sequence. What is the corresponding protein sequence?

5'-ATGGCCTGCGTTCTCAAG-3' 

PRACTICE: Even when the sequence of nucleotides for a gene is available and genomic analyses can be performed, direct chemical techniques on the physical protein are still required to determine:

- a) The molecular weight of a simple protein.
- b) The N-terminal amino acid residue.
- c) The total number of amino acid residues in the protein.
- d) The location of disulfide bonds.