

PROTEIN SYNTHESIS

①

Proteins are built up from about 20 types of amino acids. These amino acids are linked up together by peptide bond forming a long chain known as Polypeptides. The sequence of amino acids are definite for a particular protein.

The basic mechanism of protein synthesis is that DNA makes RNA, which in turns makes protein.

The central dogma of protein synthesis is as:

DNA $\xrightarrow{\text{Replication}}$ DNA $\xrightarrow{\text{Transcription}}$ RNA $\xrightarrow{\text{Translation}}$ Protein

1. Protein synthesis is broadly constitutes of two aspects.
 - I \rightarrow Transcription.
 - II \rightarrow Translation.

I \rightarrow Transcription - DNA strand ~~and~~ unwinds.

One of the two strands form an m-RNA. The nucleotides of m-RNA are complementary to template DNA ~~strand~~ strand. Uracil (U) replaces Thymine (T) of DNA. Thus A-U, G-C. Complementation is formed. After complete formation m-RNA passes to cytoplasm through nuclear pore.

Transcription needs \rightarrow template (Normally a DNA).

\rightarrow Ribonucleoside triphosphates (ATP, GTP, UTP & CTP).

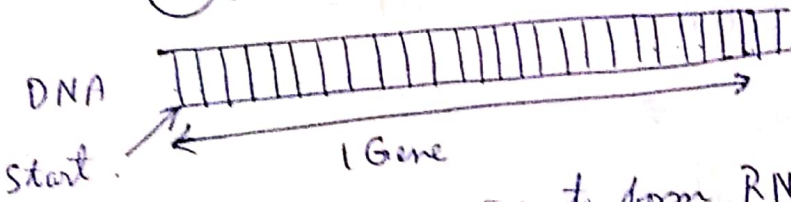
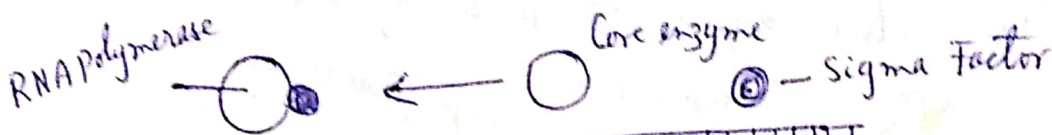
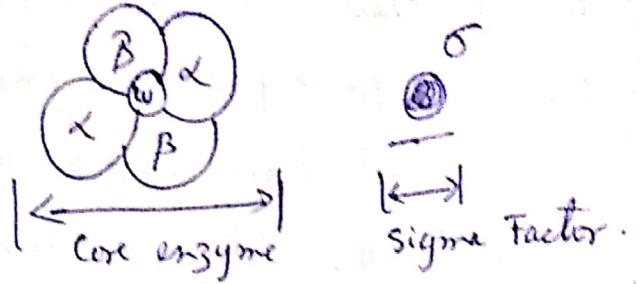
\rightarrow The enzyme RNA polymerase and

\rightarrow Divalent metal ions.

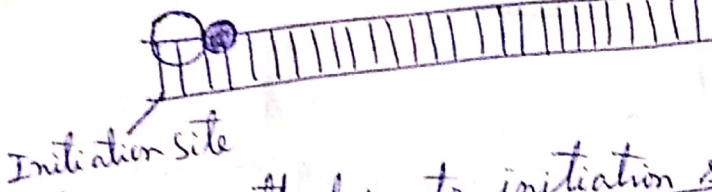
- RNA polymerase consists of a core enzyme, with sub-units $\alpha, \alpha, \beta, \beta, \omega$ and a sigma (σ) factor.

- This sigma (σ) factor initiates transcription of m-RNA on DNA template and the core enzyme constitute transcription.

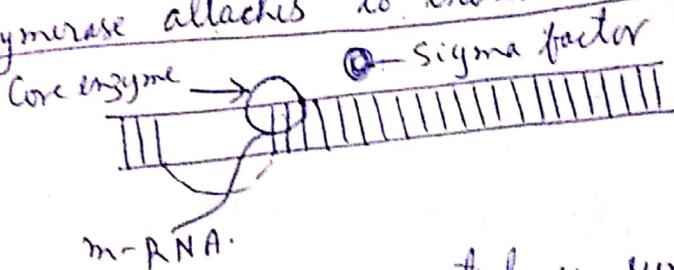
- All RNA chains start with PPP G or PPP A
- They are synthesized in 5' → 3' direction.
- rho (ρ) factor is required for termination of mRNA-RNA synthesis.



[A] Sigma factor and core enzyme join to form RNA polymerase



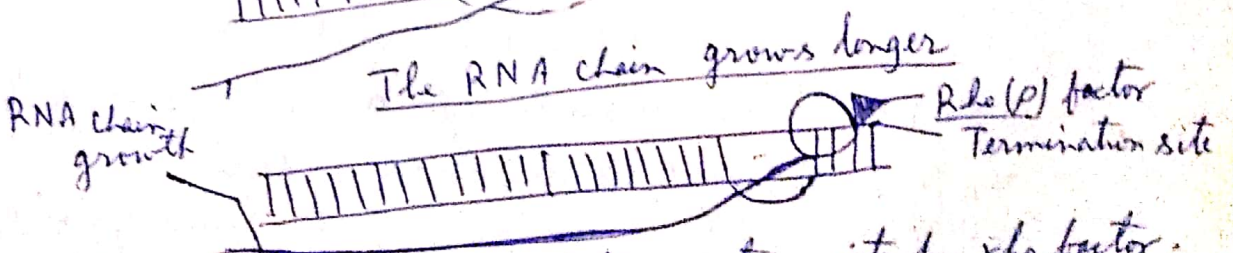
[B] RNA polymerase attaches to initiation site.



[C] DNA unwinds and core enzyme catalyses synthesis of m-RNA. Sigma factor dissociates from core enzyme.



[D]



[E] Termination of chain growth at termination site by rho factor.

II TRANSLATION:

The m-RNA has a series of triplet bases, known as codons, which pair with triplets of anticodons on t-RNA.

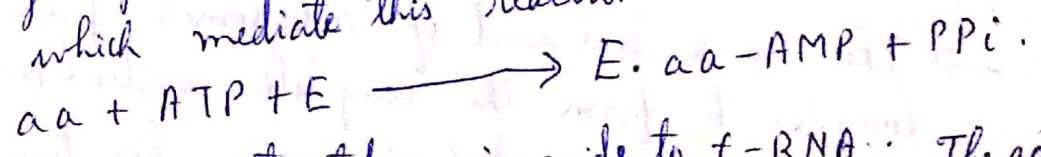
Triplets of m-RNA are according to a series of bases on DNA. Triplets of t-RNA arranged come in contact ~~also~~ in accordance with triplets of m-RNA. Hence series of amino acids are according to m-RNA and DNA in turn. So it may be said that proteins are according to amino acid chains or m-RNA or DNA.

~~The genetic information present on m-RNA~~

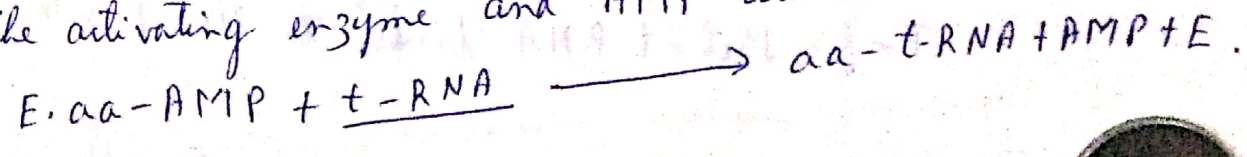
The process of translation involves:-

- (a) Activation of amino acids.
- (b) Transfer of activated amino acid to t-RNA.
- (c) Initiation of polypeptide chain synthesis.
- (d) Chain elongation
- (e) Chain termination.

(a) Activation of amino acids:- The amino acids (aa) used in protein synthesis are activated by ATP. ^{forming} Aminoacyl adenylates (aaa) or aminoacyl AMP_a is formed. In this reaction pyrophosphates (PPi) are released. ~~The enzyme~~ ~~and~~ ~~mediate~~ this reaction. Aminoacyl synthetase (E) ~~acti~~ is the specific activating enzyme which mediate this reaction.



(b) Transfer of activated amino acids to t-RNA: The activated amino acids is transferred to the specific t-RNA. The activating enzyme and AMP are released.



(c) Initiation of Chain Synthesis:-

This process requires:

③ decider

- i) sub-units of ribosomes.
- ii) m-RNA.
- iii) a source of energy GTP.
- iv) activated amino acids attached to t-RNA.
- v) Initiation factor (IF).

→ In prokaryotes these initiation factors are -
IF-1; IF-2 and IF-3.

→ In eukaryotes these initiation factors are -
eIF-2; eIF-2'; eIF-2a; eIF-2a₂
eIF-2a₃ and eIF-3.

This process goes on as follows:

→ The 30S sub unit of ribosome attaches to m-RNA to form an m-RNA-30S complex, with the help of IF-3 and Mg^{++} .

→ Initiation codon is generally AUG, which ^{attaches to} ~~is for~~ anticodon UAC on t-RNA forming 30S initiation complex, ~~then~~ with the help of initiation factors IF-2; IF-3 and GTP as source of energy.

- The first amino acid ~~is~~ is N-formyl methionine (f-met) in prokaryotes and methionine in eukaryotes.

→ The 50S sub-unit of ribosomes joins to 30S initiation complex to form the complete initiation complex (70S).

→ The larger sub-unit of ribosomes has two binding sites for t-RNA —

- x) A-site or Acceptor site and
- x) P-site or Peptidyl site

f-Met-tRNA binds to the P site.