

(d) Chain Elongation: The elongation of polypeptide chain needs certain factors, known as Elongation Factor (EF). (5)

~~These are EF-Tu, In prokaryotes - EF-1~~

In prokaryotes these are \rightarrow EF-Tu; and EF-Ts and EF-G
and in Eukaryotes \rightarrow EF-1 and EF-2.

The process goes on as \rightarrow

(i) The second amino acid-t-RNA complex (aa_2 -tRNA) comes to A-site of larger sub-unit. It needs EF-Tu, GTP and Mg^{++}

(ii) Peptide bond is formed by transfer of f-Met to the second amino acid (aa_2); ^{It needs} Mg^{++} and K^+ in addition to the enzyme peptidyl transferase.

(iii) Translocation \rightarrow This aa_2 -tRNA complex moves from the A-site to the P-site.
It needs EF-G, GTP and Mg^{++} .

(iii) There is movement of ribosome related to mRNA in $5' \rightarrow 3'$ direction.

The tRNA which carried f-Met get released from f-Met and come down from the position 'P' releasing the ribosome.

The third amino acid-t-RNA complex (aa_3 -tRNA) now come to A-site which has been vacated by second amino acid-t-RNA (aa_2 -tRNA) complex.

(iv) In this way elongation of chain continues till the onset of termination codons UAA, UAG or UGA on mRNA.

(e) Chain Termination :- This process requires

— A termination codon on the mRNA i.e.

UAA

UAG

UGA

— Release factors

x) In prokaryotes RF-1; RF-2; and RF-3.

x) In eukaryotes RF.

In prokaryotes Enzyme Deformylase and Aminopeptidase and in Eukaryotes only Aminopeptidase
The process occurs as follows.

i) For attachment of release factor to ribosomes, the signal is provided by termination codon.

ii) These release factors interact with peptidyl transferase and cause hydrolysis of the bond between polypeptide chain and t-RNA.

(iii) Hence the ~~rest~~ chain is released from the ribosome.

(iv) ~~Hydrolysis of GTP results causes the dissociation~~

(iv) ~~Due to~~ Due to hydrolysis of GTP release factors dissociate from ribosomes.

(v) The t-RNA get released from peptide/amino acyl bonds and become unloaded.

(vi) There is dissociation of ribosomal sub-units

(vii) m-RNA is released ~~and~~ for breakdown to nucleotides.

~~(viii)~~

b) Processing of polypeptide chain: After release of polypeptide chain from ribosome, there may be cleavage of formyl residue or of methionine.

In some cases internal cleavage cuts the polypeptide chain into two or more pieces.

~~Formyl~~

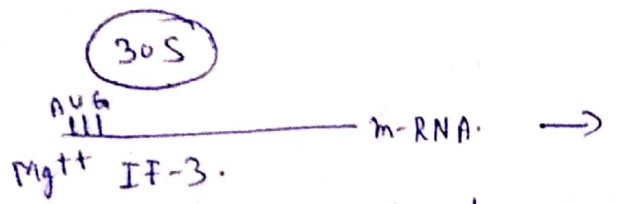
(1) Formyl methionyl

(1) Formyl methionyl peptide \longrightarrow Formic Acid + Methionyl peptide

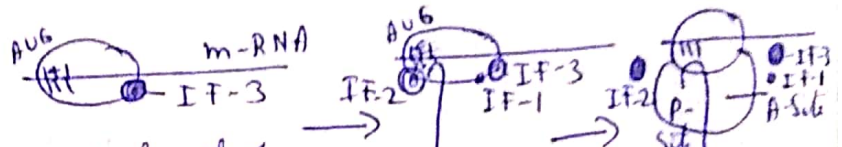
(2) Methionyl peptide \longrightarrow Methionine + Peptide.

Thus a chain may begin with alanine, serine or the threonine.

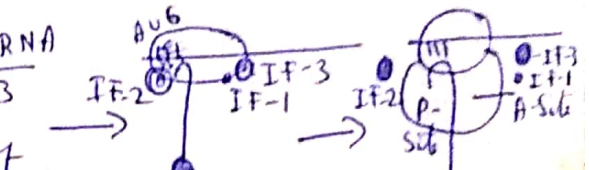
In this way a particular protein is synthesized.



A) The 30-s ribosomal Unit & m-RNA



(B) Attachment of m-RNA to 30-s sub-unit, the process needs IF-3, & Mg⁺⁺

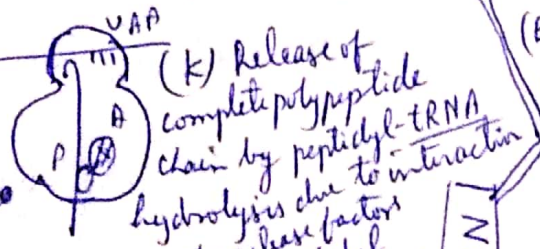
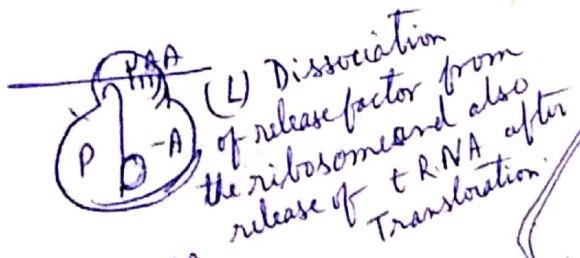


(C) Attachment of f-Met-tRNA to initiation codon AUG of m-RNA to form 30-s initiation complex. It needs IF-2, IF-1, & IF GTP.

(D) Joining of 50-s to 30-s to form initiation complex.

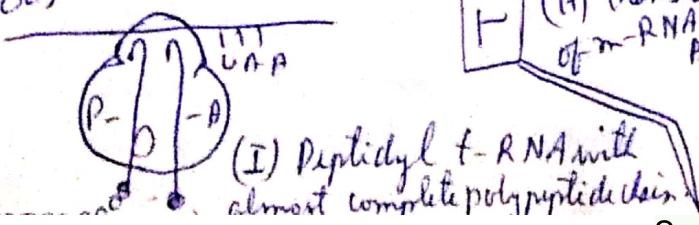
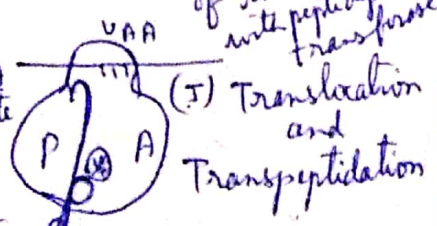
Initiation

m-RNA (30S) (50S) (M)
Dissociation of Ribosomes and release of mRNA



TERMINATION

Termination codon UAA comes to recognition site. Polypeptide chain is attached to t-RNA. Release Factor interacts with ribosomes in response to the termination codon.



TRANSLATION

