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TOPIC: Translation

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Introduction to Translation

- Translation: The biosynthesis of a protein or a polypeptide inside a living cell.
- In process of translation the language of nucleotides sequence on mRNA is translated in to the language of amino acid sequence.
- It occur in cytoplasm where ribosomes are located
- It is a universal process

- In translation ,messenger RNA is decoded to produce a specific polypeptide
- This uses mRNA sequence as a template to guide the synthesis of a chain of amino acid that form protein
- Many types of transcribed RNA, such tRNA,rRNA,snRNA are not necessarily translated to amino acid sequence

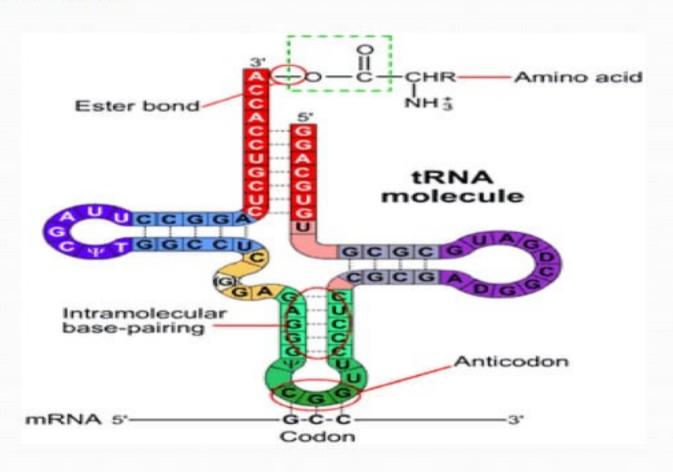
Steps of Translation

- Translation proceeds in four phases
 - Activation
 - Initiation
 - Elongation and
 - Termination

Most studies do not include activation as a step of translation

Activation

- In activation, the correct amino acid is covalently bonded to the correct tRNA
- While technically this not a step in translation, it is required for translation to proceed
- The amino acid is joined by its carboxyl group to the 3`
 OH of tRNA by an ester bond with help of ATP
- When tRNA has an amino acid linked to it, it is termed as "charged"



Initiation

- Prokaryotes initiation require the large and small subunits, the mRNA, the initiator tRNA and three initiation factors (IF-1, IF-2, IF-3) and GTP.
- IF-3 binds to the free 30s subunit, this help to prevent large subunit binding to it without mRNA and forming an inactive ribosome
- IF-2 complexed with GTP and IF-1 binds to small subunit. It will assist the charged initiator tRNA to bind

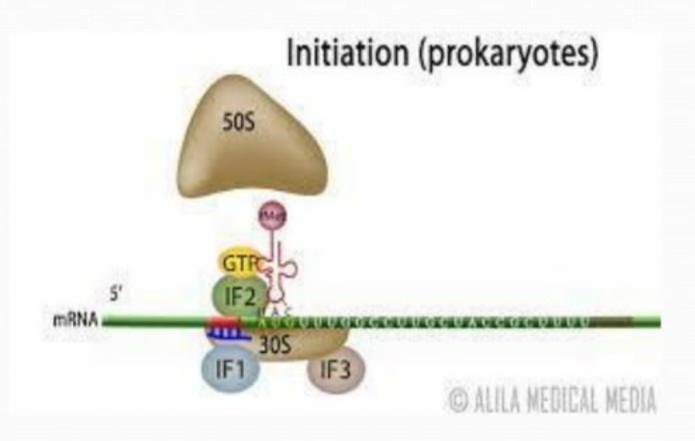
- The 30s subunit attached to a mRNA molecule making use of the ribosomal binding site on mRNA
- The initiator tRNA can then bind to the complex by base pairing of its anticodon with AUG codon on mRNA
- At this point,IF3 can be released, as its role in keeping the subunits apart are complete
- This complex is called <u>30s initiation complex</u>

 The 50s subunit can now bind, which displace IF1 and IF2, and the GTP is hydrolysed in this energy consuming step

This complex is called <u>70s initiation complex</u>

- The assembled ribosome has two tRNA binding sites
- These are called the A and P sites, for amino acyl and peptidyl sites and one site is E (exit site) for empty tRNA
- The A site is where incoming amino acyl tRNA molecule bind, and the p site is where the growing polypeptide chain is usually found
- The sites are in cleft of small subunit and contain adjacent codon that are being translated

- One major of initiation is the placement of initiator tRNA in the P site
- It is the only tRNA that does this, as all other must enter the A site



Elongation

- With the formation of 70s initiation complex the elongation cycle can begin
- It involves three elongation factors EF-Tu, EF-Ts and EF-G, GTP, charged tRNA and the 70s initiation complex

Elongation is divided into 3 steps

Amino acyl tRNA delivery.

- EF-TU is required to deliver the amino acyl tRNA to A site and energy is consumed in this step by hydrolysis of GTP
- The released EF-Tu GDP complex is regenerated with the help of EF-TS
- In the EF-Tu EF-Ts exchange cycle EF-Ts displaces the GDP and replace itself by GTP
- The resultant EF-Tu.GTP complex is now available to bind another amino acyl tRNA and deliver it to ribosome

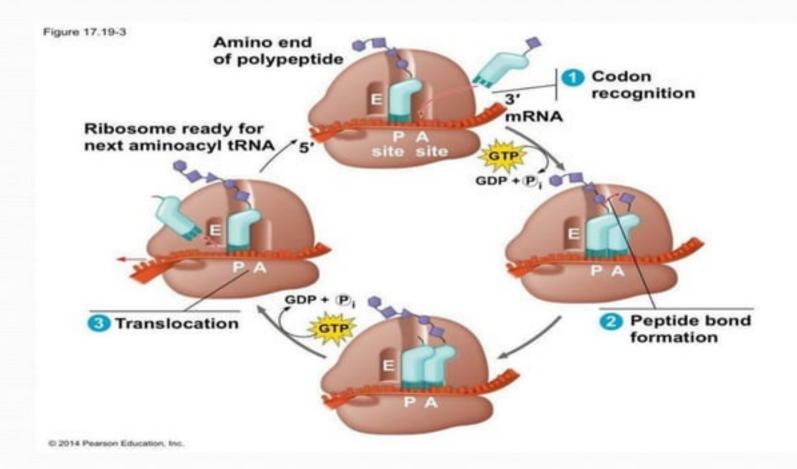
 All amino acyl tRNA can form this complex with EF-Tu except the initiator tRNA

2 .Peptide bond formation.

- After aminoacyl-tRNA delivery, the A and P sites are both occupied and the two amino acids that are to be joined are close to each other
- The peptidyl transferase activity of the 50s subunit can now form a peptide bond between the two amino acids

3. Translocation

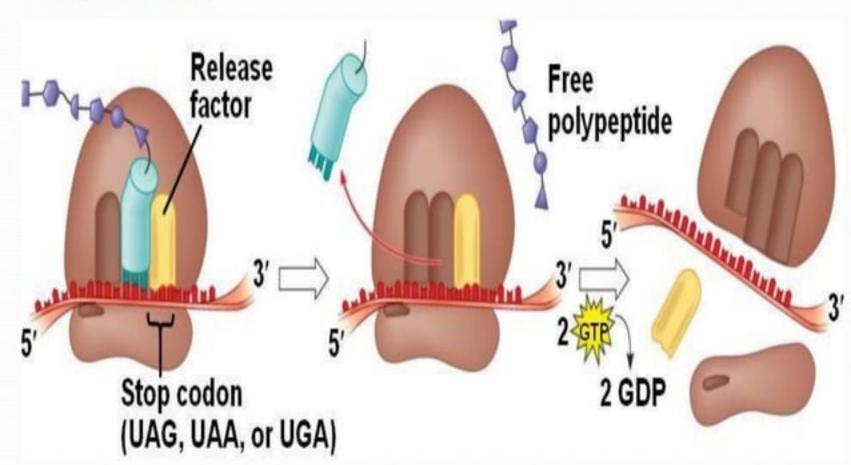
- A complex of EF-G(translocase) and GTP binds to the ribosome and ,is an energy consuming step, the discharged tRNA is ejected from the P site, the peptidyltRNA is moved from A site to P site
- The mRNA moves by one codon relative to one codon to the ribosome
- GDP and EF-G are released . A new codon is now present in the vacant site



Termination

- Termination of translation happens when the A site of the ribosome faces a stop codon (UUA,UGA or UGA)
- When this happens, no tRNA can recognize it, but a releasing factor can recognize the stop codons and causes the release of polypeptide chain
- In prokaryotes once a stop codon occupies the A site, three termination or release factor (RF1,RF2,RF3) contribute to the hydrolysis of peptidyl-tRNA bond

- Release the free polypeptide and last uncharged tRNA from P site
- The dissociation of the 70 s into 30s and 50s subunits
- RF1 binds A site and release the polypeptide and uncharged tRNA
- RF2 releases the RF1 from A site and release itself as well from translation binding site(present on large subunit)
- RF3 function unknown
- Another factor called Ribosomal releasing factor causes the dissociation of 70s complex



Translation in Eukaryotes

Eukaryotic Translation

- In prokaryotic cell, transcription and translation are coupled, that is, translation begins while the mRNA is still being synthesized. In a eukaryotic cell, transcription occur in the nucleus, and translation occur in the cytoplasm.
- Translation process in eukaryotes involve
 - Activation (not essentially the step of translation. This occur the same way as in prokaryotes)
 - Initiation
 - Elongation and
 - termination

1. Initiation

 The initiation of translation in eukaryotes is complex, involving at least 10 initiation factors (eIFs) and is divided into three steps:

- a) Formation of 43s preinitiation complex.
- Formation of 48s initiation complex.
- Formation of 8os initiation complex.

a. Formation of 43s preinitiation complex

- A ternary complex containing met-tRNA and eIF-2 bound to GTP attaches to 40s ribosomal subunit to form 43s preinitiation complex.
- The presence of eIF-3 and eIF-1A stabilizes this complex.

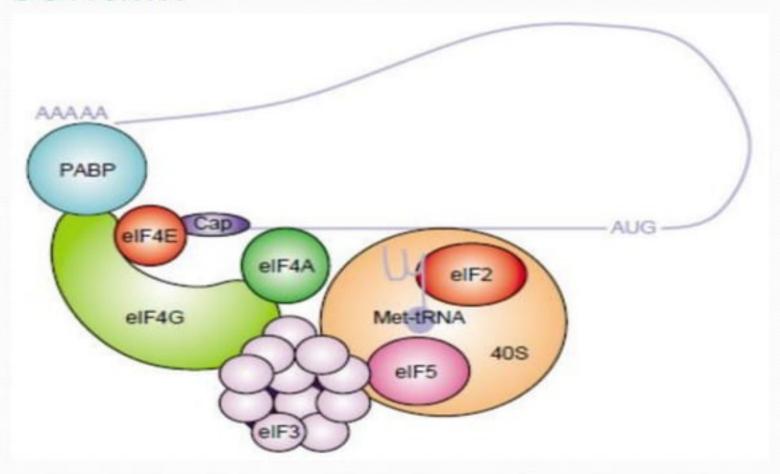
b.Formation of 48s initiation complex

- The binding of mRNA to 43s preinitiation complex results in formation of 48s initiation complex.
- eIF-4f is formed by the association of eIF-4G, eIF-4A with eIF-4E
- The eIF-4F(referred to as cap binding protein) binds to the cap of mRNA.

- Then eIF-4A and eIF-B binds to mRNA and reduces its complex structure.
- This mRNA is then transferred to 43s complex
- The ribosomal initiation complex scans the mRNA for identification of appropriate initiation codon
- 5'-AUG is the initiation codon

c.Formation of 80s initiation complex

- 48s initiation complex binds to 6os ribosomal subunit to form 8os initiation complex
- The binding involves hydrolysis of GTP(bound to eIF-2)
- This step is facilitated by the involvement of eIF-5
- As the 8os complex is formed, the initiation factors bound to 43s initiation complex are released and recycled



2. Elongation

- Ribosomes elongate the polypeptide chain by sequential addition of amino acids
- The amino acid sequence is determined by the order of the codons in the specific mRNA
- Elongation, a cyclic process involving certain elongation factors(EFs)
- Elongation may be divided into three steps
- Binding of aminoacyl-tRNA to A-site
- Peptide bond formation
- translocation

a.Binding of Aminoacyl t-RNA to Asite

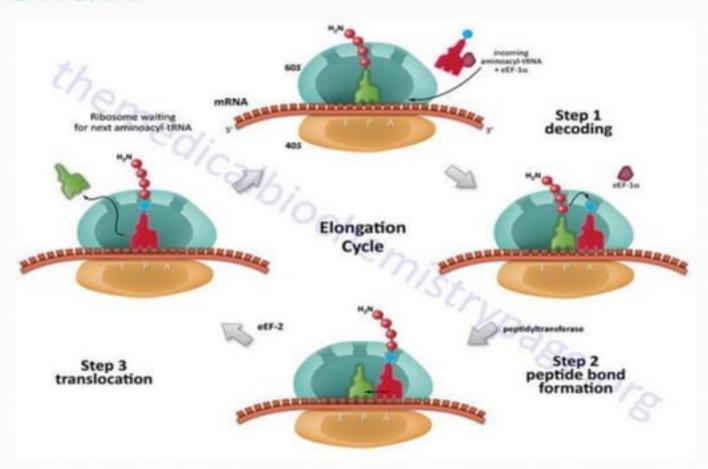
- The 8os initiation complex contains met tRNA in the P-site and A- site is free
- Another aminoacyl-tRNA is placed in the A site
- This require proper codon recognition on mRNA and involvement of EF-1a and supply of energy by GTP
- The aminoacyl –tRNA is placed in the A-site, EF-1a and GDP are recycled to bring another aminoacyltRNA

b. Peptide bond formation

- The peptidyl transferase catalyzes the formation of peptide bond
- Net result of peptide bond formation is the attachment off the growing peptide chain to the tRNA in A-site

c. Translocation

- The ribosome moves to the next codon of mRNA (towards 3' end)
- This process is called translocation, involves the movement of growing peptide chain from A-site to Psite
- Translocation require EF-2 and GTP
- GTP get hydrolyzed and supplies energy to move mRNA
- EF-2 and GTP complex recycles for translocation

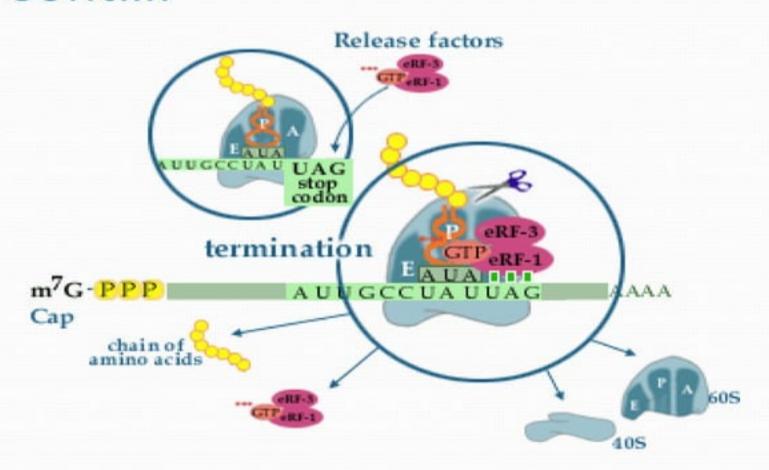


3. Termination

- One of the stop signals (UGA,UAA and UGA) terminates the growing polypeptide
- When the ribosome encounter s a stop codon, there is no tRNA available to bind to the A site of ribosome
- Instead a release factor binds to it
- In eukaryotes eRF1 recognizes all the three stop codons, and eRF3 stimulate the termination events

Cont...

- Once the release factor binds, the ribosome unit falls apart
 - releasing the large and small subunits
- the tRNA carrying the polypeptide is also released, freeing up the polypeptide product.



References

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Thank you