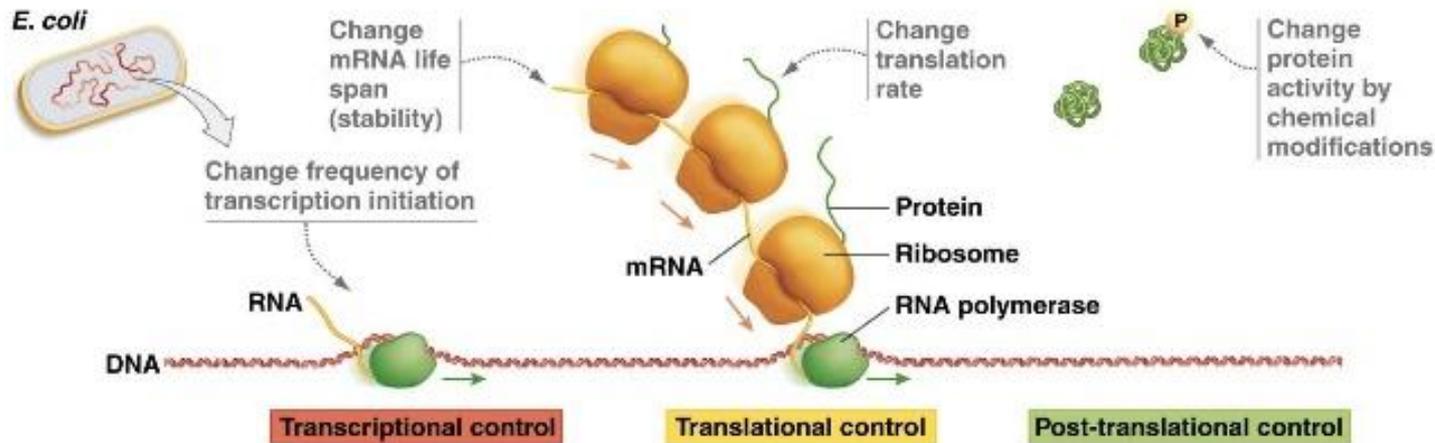


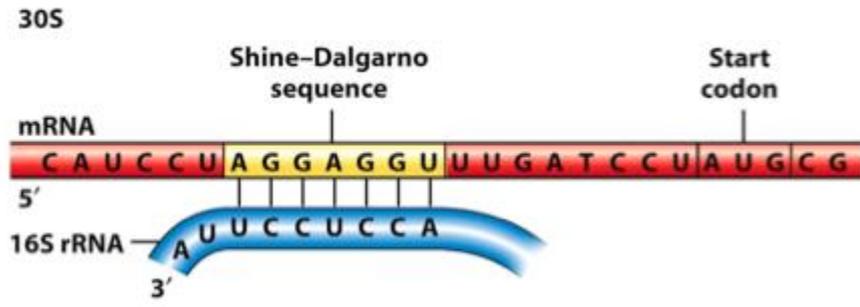
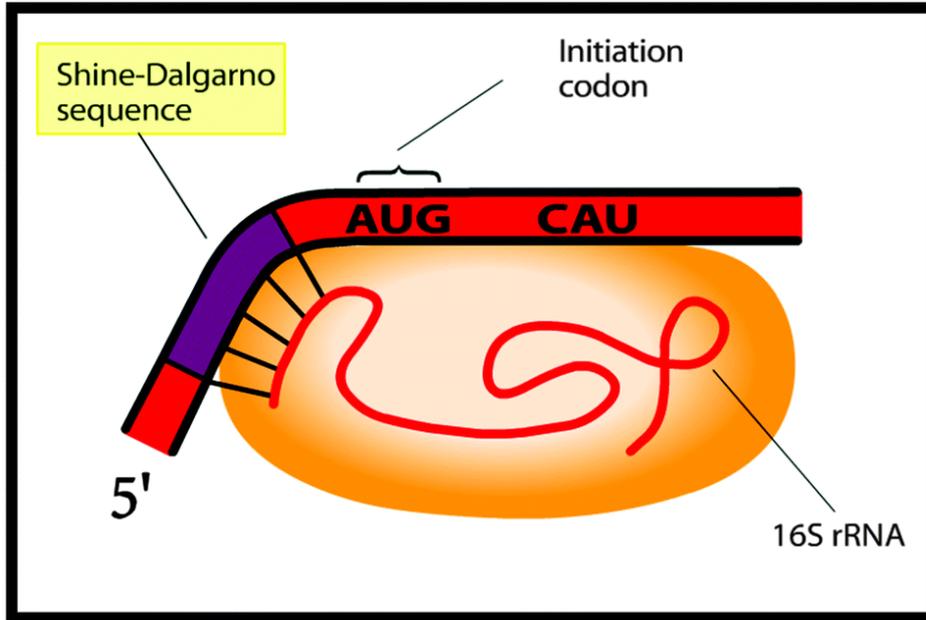
# Regulation of Gene Expression



- Gene expression can be regulated:
  - During transcription (transcriptional control).
  - During translation (translational control).
  - After translation (post-translational control).

# Translational control in prokaryotes

- In prokaryotes translation begins with binding of ribosome to a specific sequence in the messenger RNA - Shine-Dalgarno (SD) Sequence. SD is a ribosomal binding site generally located around 8 bases upstream of the start codon AUG. The six-base consensus sequence is AGGAGG. It is complementary to a specific region of 16S ribosomal RNA.

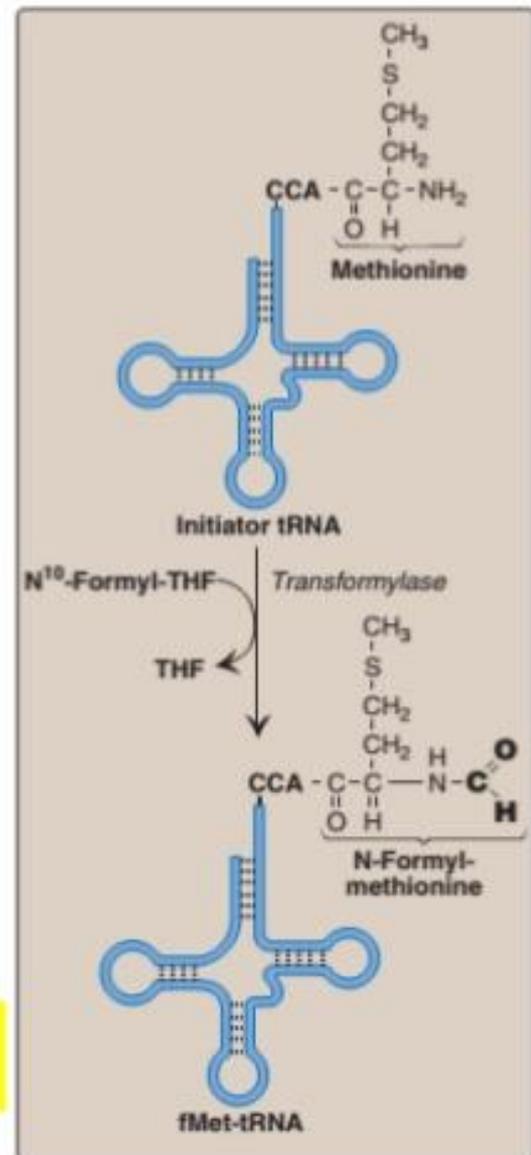


# Initiation codon

25

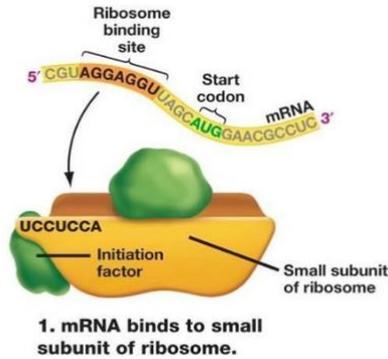
- Initiating “AUG” is recognized by special initiator tRNA.
- Recognition is facilitated by **IF-2 (bound to GTP) in Prokaryotes** and **eIF2-GTP in Eukaryotes**.
- The AA charged initiator tRNA enters the ribosomal P site, and GTP is hydrolysed to GDP.

**NOTE:** The initiator tRNA is the only tRNA recognized by eIF-2 and the only tRNA to go directly to the P site.



## Translation initiation in bacteria

### Shine-Dalgarno sequence

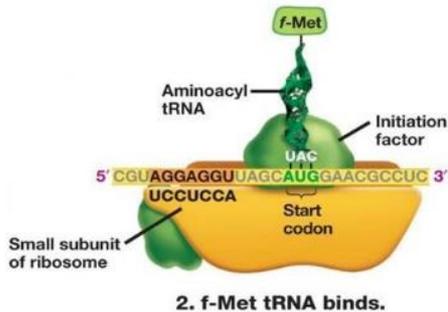


*m*

## Translation initiation in bacteria

### Initiator tRNA in bacteria

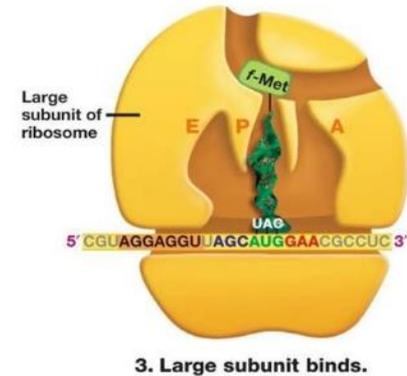
1. The initiator tRNA (fMet-tRNA) gets carried to the complex (30S ribosome + IF1 + IF3) by initiation factor IF2 using GTP.



*m*

## Translation initiation in bacteria

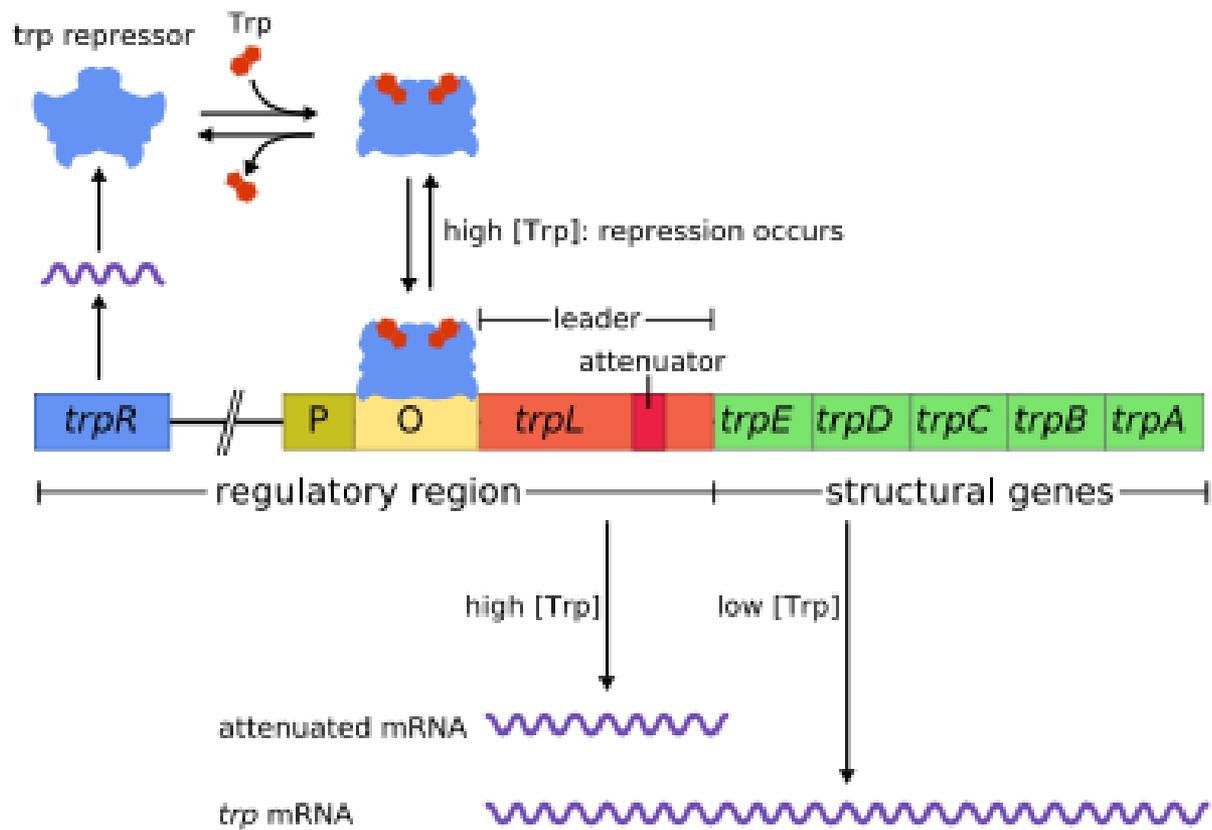
- The initiation factors (IF1 and IF3) gets released and the resulting complex is called **the initiation complex**.



*m*

# Attenuation of trp operon

- Attenuation is a second mechanism of negative feedback in the trp operon. The repression system targets the intracellular trp concentration whereas the attenuation responds to the concentration of charged tRNA<sup>trp</sup>
- Attenuation results in only 10% transcription rate of the trp operon structural genes
- Attenuation is made possible by the fact that **in prokaryotes** (which have no nucleus), **the ribosomes begin translating the mRNA while RNA polymerase is still transcribing the DNA sequence**. This allows the process of translation to affect transcription of the operon directly.

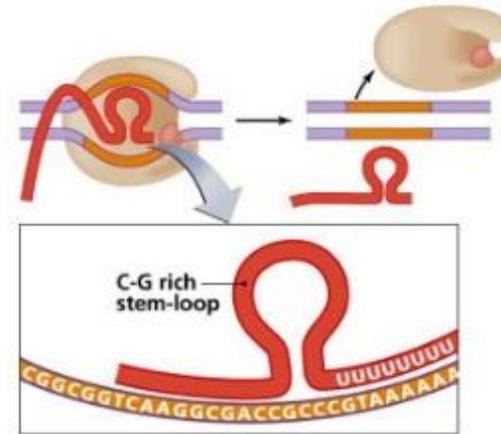


# Bacterial transcription termination

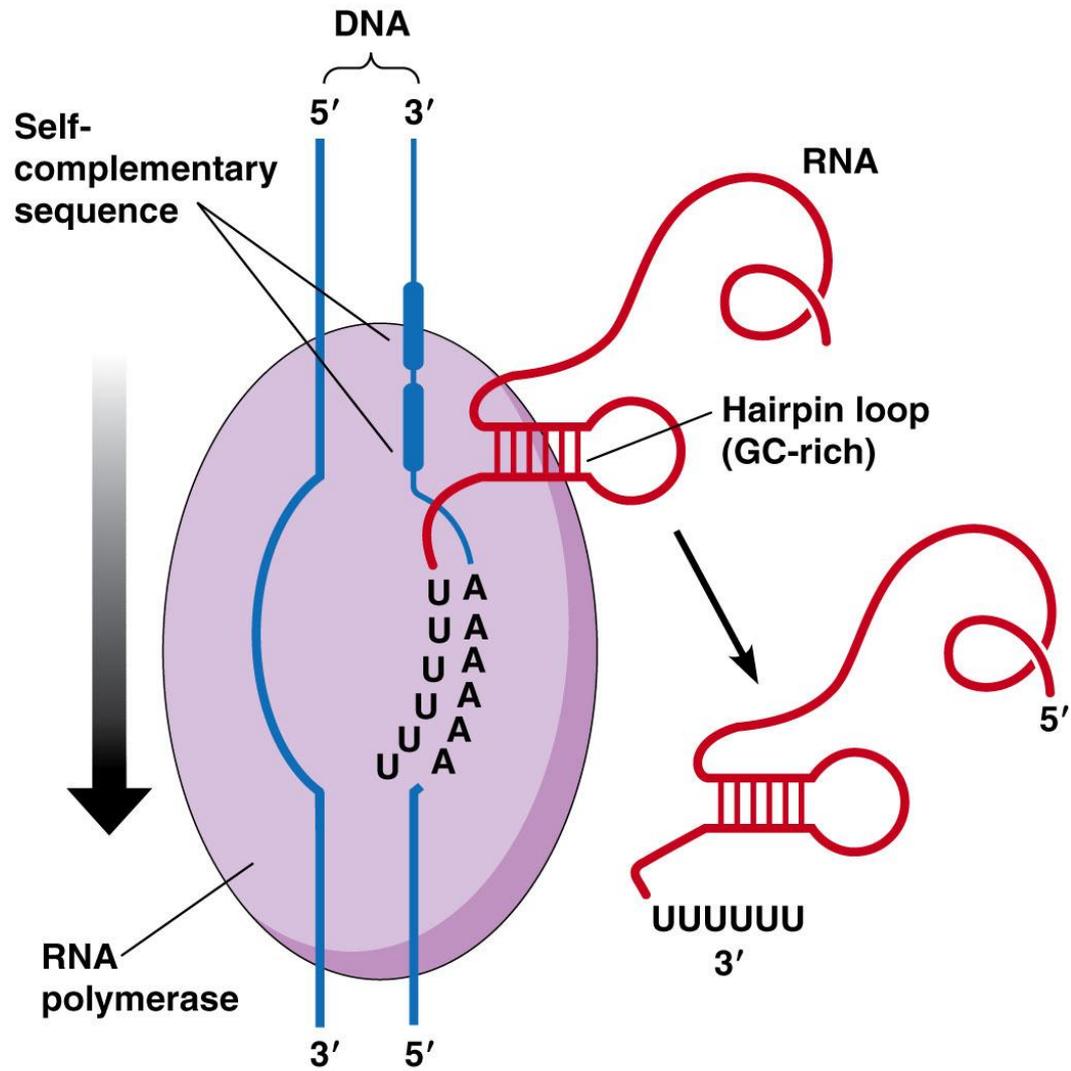


## Termination (Rho-independent terminator) - type 1 terminator

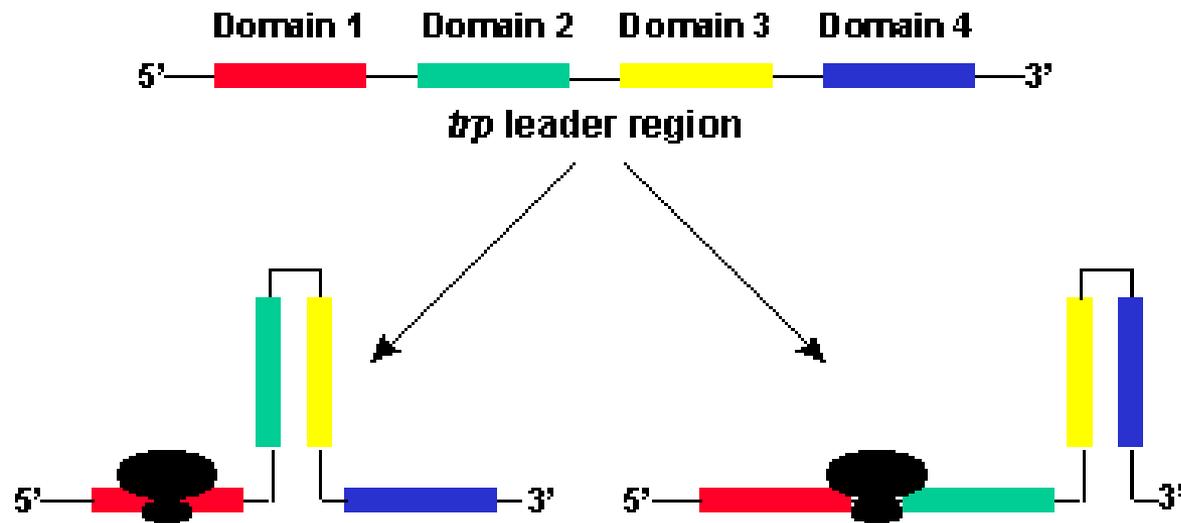
- RNA moves past the inverted repeats and transcribes the termination sequence.
- Because of the inverted repeat arrangement → RNA synthesized forms a hairpin loop structure.
- Hairpin loop makes the RNA polymerase slow down and eventually stops.



(c) Termination of transcription



## Attenuation of the *trp* operon mRNA



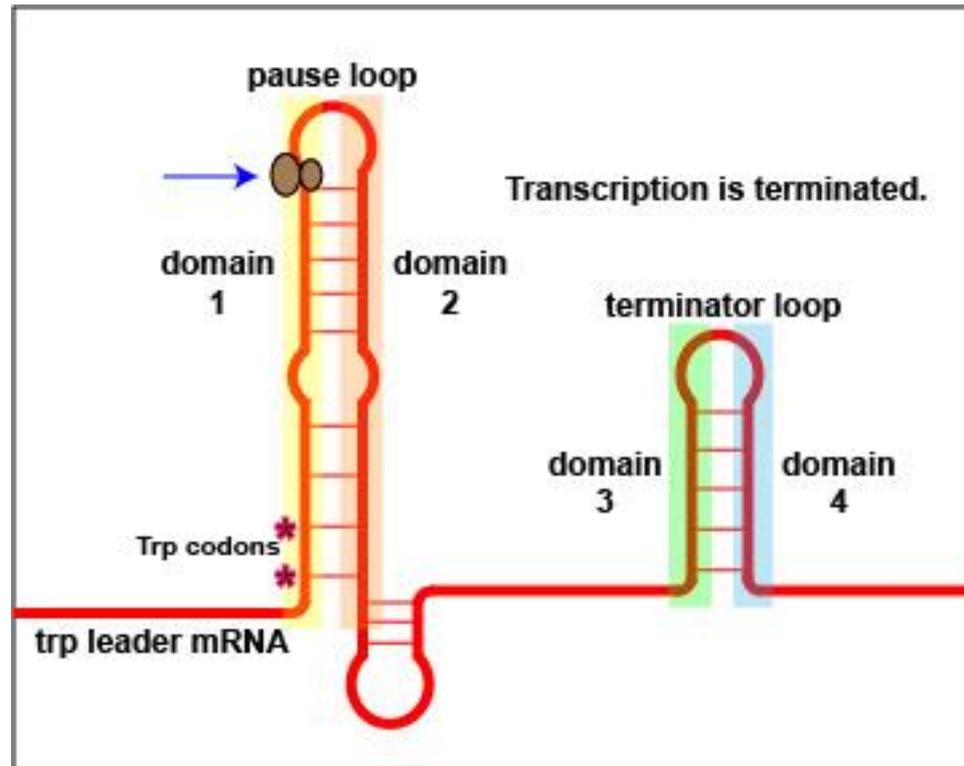
### Low Tryptophan Levels

- Slow translation of Domain 1 peptide
- Domain 2-3 pairing occurs
- Normal full gene transcription

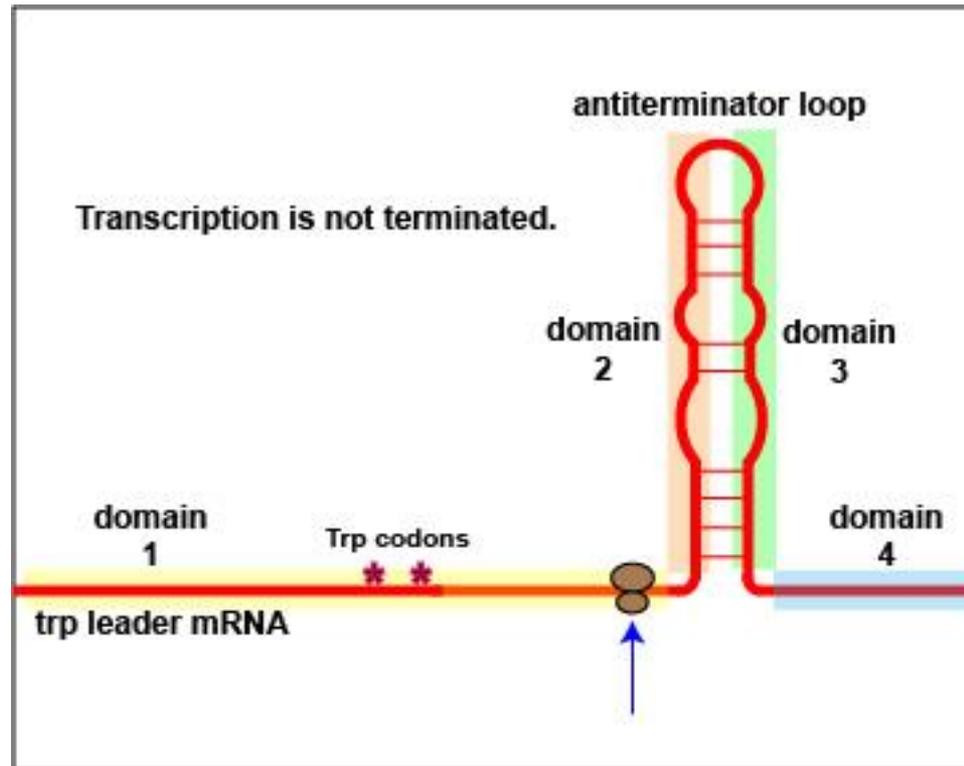
### High Tryptophan Levels

- Fast translation of domain 1 peptide
- Domain 2 blocked by ribosome
- Domain 3-4 pairing occurs
- Attenuation of transcription occurs
- Only 10% of normal mRNAs made

# Attenuation, tryptophan present



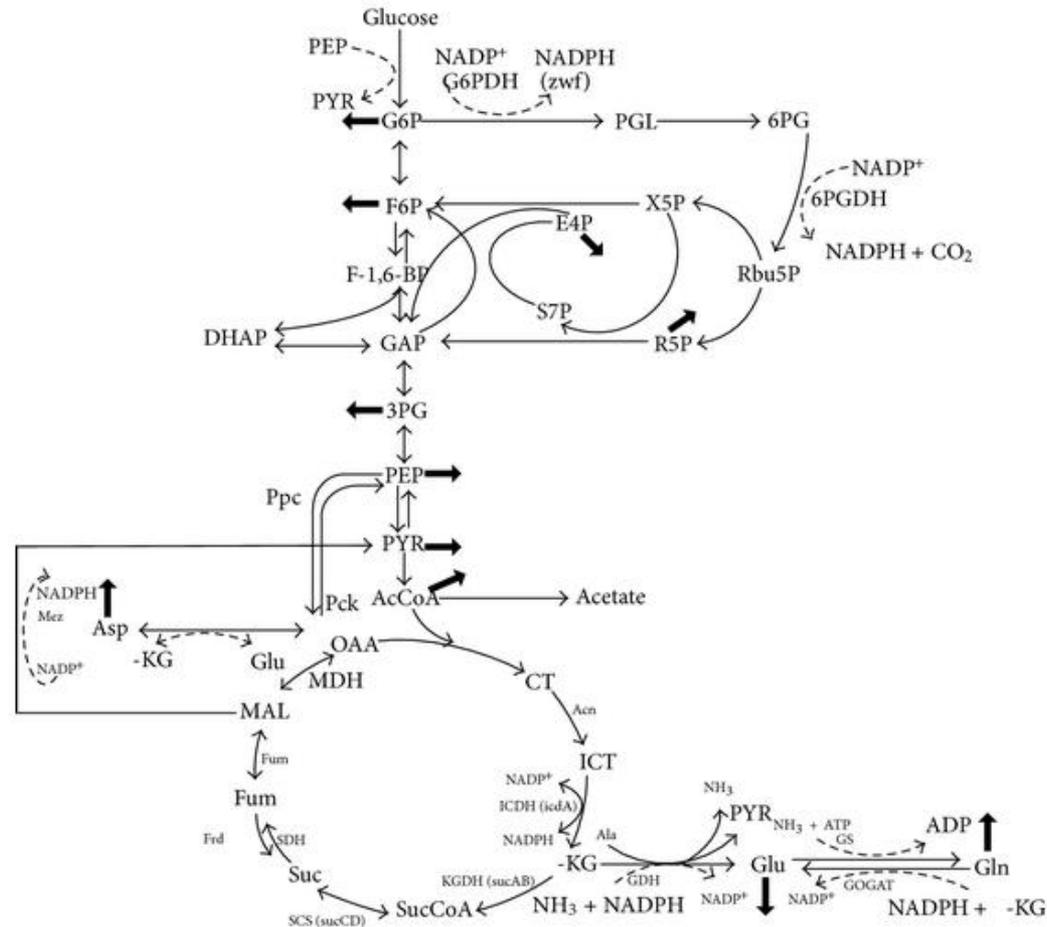
# Attenuation, no tryptophan



# Biochemical Pathways

- A biochemical pathway (also called a metabolic pathway) is a series of enzyme-mediated reactions where the product of one reaction is used as the substrate in the next. Each enzyme is coded by a different gene.
- A network of complex interlocking mechanisms regulates expression of the enzymes.

# Biochemical pathway of glucose catabolism in *E.coli*





# Unrelated fact

- The Perseverance rover landed on Mars last Tuesday. What is it looking for?



**Strelley Pool Australian Archaean Stromatolites – 3.43 billion years old**