

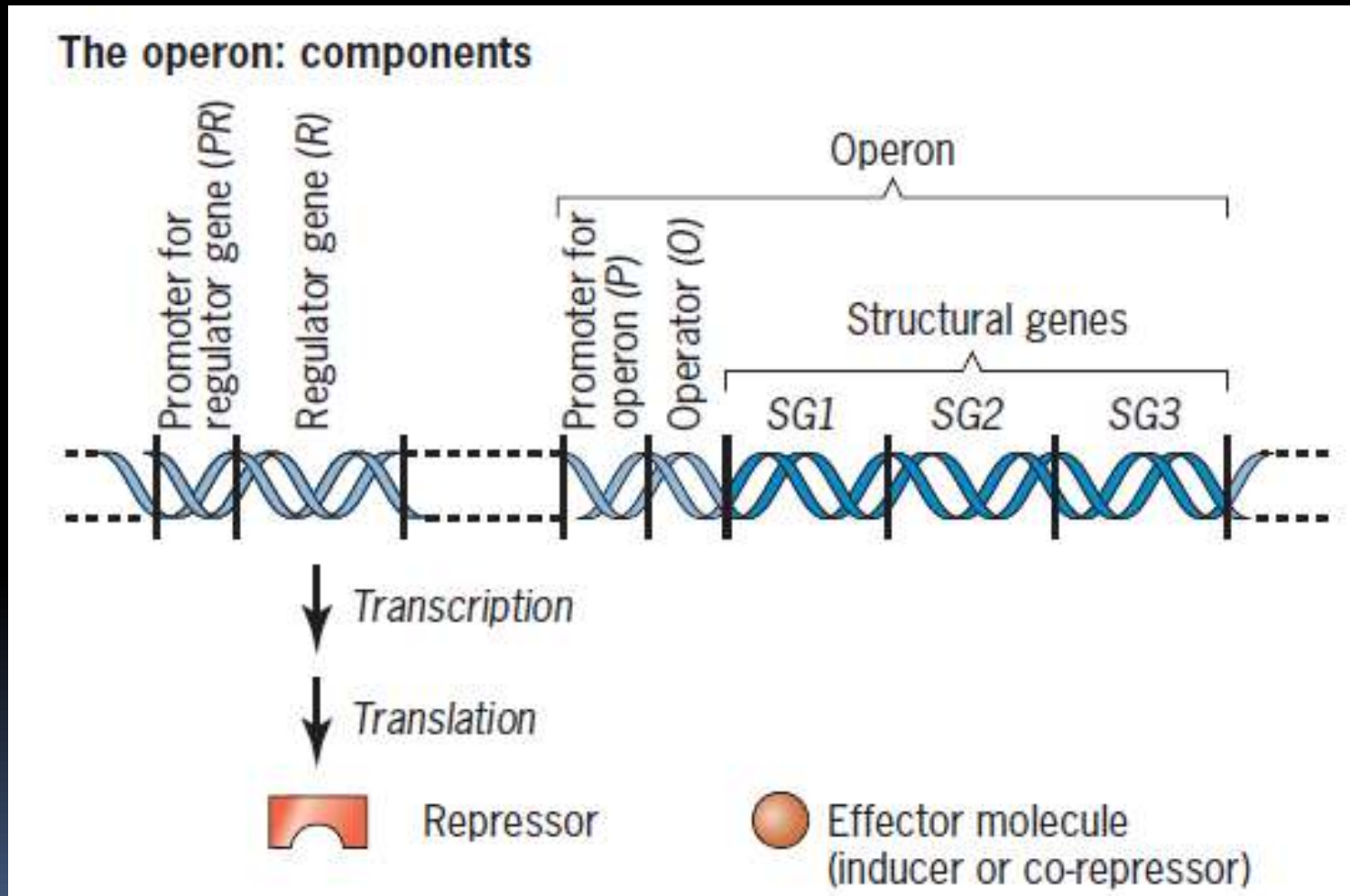
**ZOO 203 – MOLECULAR BIOLOGY**  
**UNIT 4: REGULATION OF GENE EXPRESSION IN PROKARYOTES**  
**(PART- II)**


**TRYPTOPHAN**  
**OPERON**

By

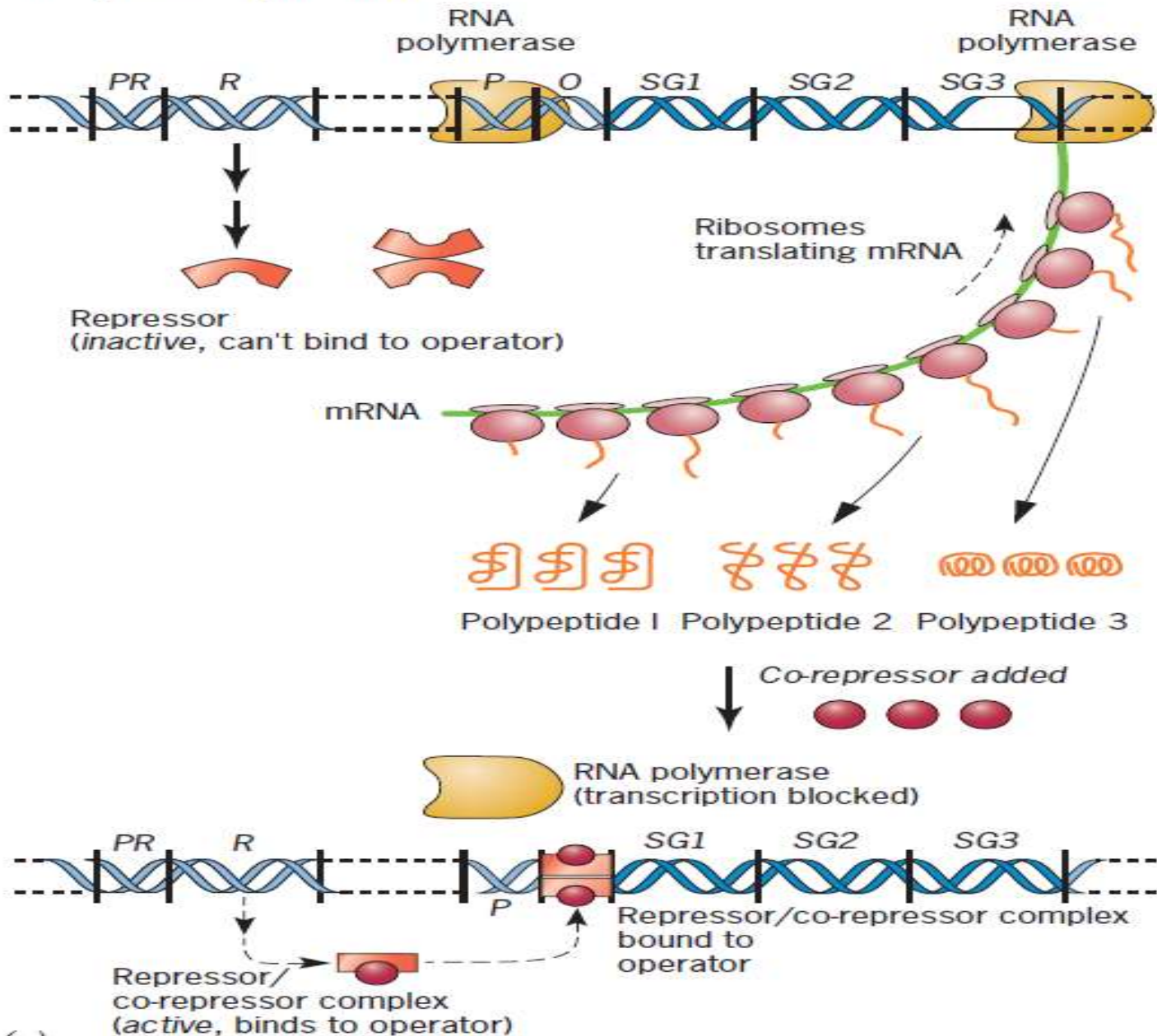
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**DEPT. OF ZOOLOGY**  
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**Components of an operon:** one or more structural genes (three, *SG1*, *SG2*, and *SG3*) and the adjoining operator (*O*) and promoter (*P*) sequences

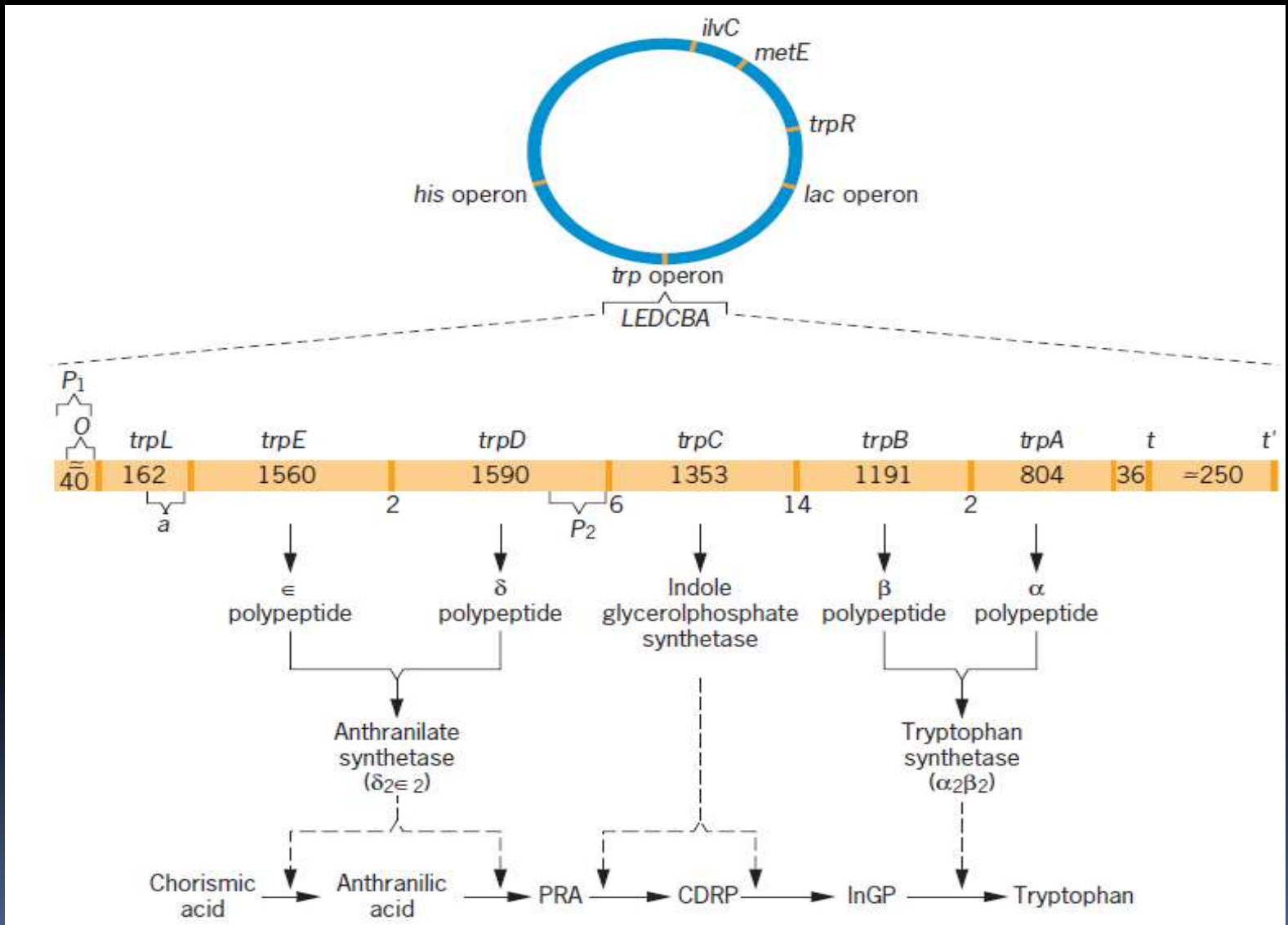


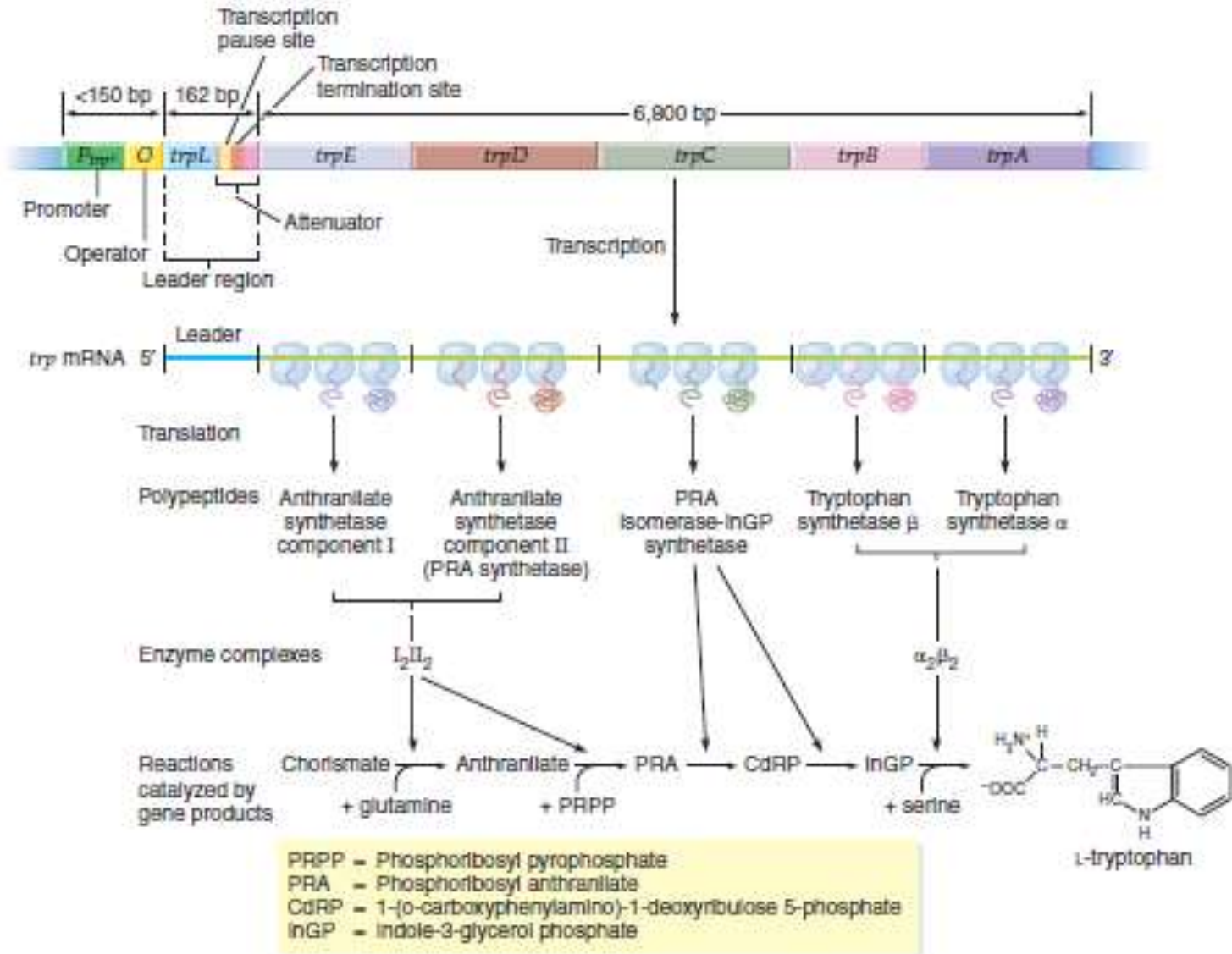
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- The *trp* operon of *E. coli* is a negative repressible operon
  - analyzed in detail by Charles Yanofsky and colleagues

# The operon: repression



# Organization of the trp (tryptophan) operon in *E. coli*.







Two regulatory mechanisms are involved:

- repressor–operator interaction
- other determines whether initiated transcripts include the structural genes or are terminated before those genes are reached




❖ **The rate of transcription of the trp operon in absence of tryptophan (derepressed state) is 70 times the rate that occurs in presence of tryptophan (repressed state)**

**In trpR mutants, which lack a functional repressor, the rate of synthesis of the tryptophan biosynthetic enzymes is still reduced about tenfold by the addition of tryptophan to the medium.**


**This additional reduction in trp operon expression is caused by attenuation**



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- Attenuation can reduce transcription of the trp operon by a factor of 8 to 10. Thus, repression and attenuation together can regulate the transcription of the trp operon by a factor of about 560 to 700.

# ATTENUATION

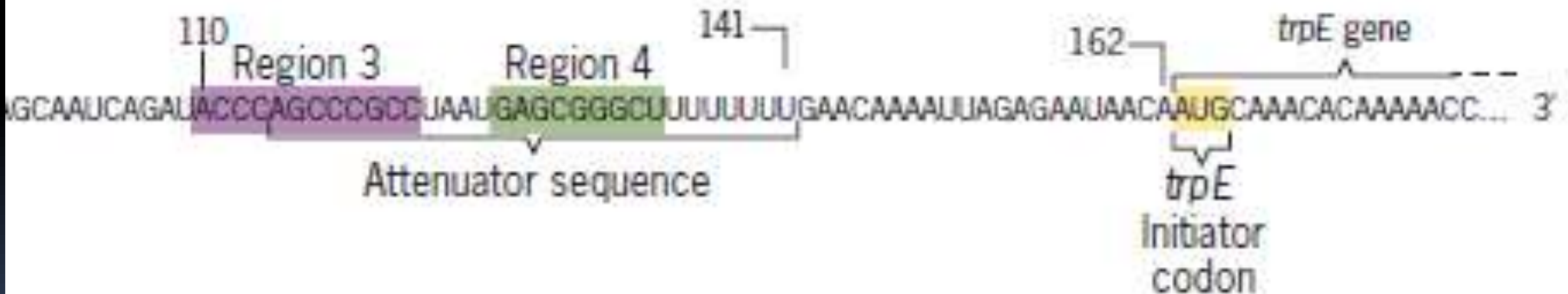
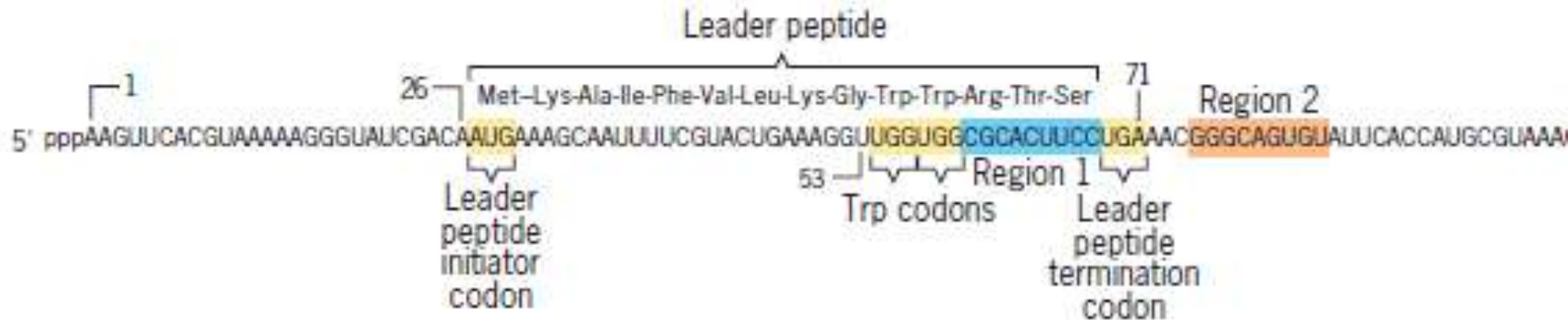
- The regulation of the trp operon other than repression is called **attenuation**
- the sequence within trpL that controls this phenomenon is called the **attenuator**
- Attenuation controls the termination of transcription at a site near the end of the mRNA leader sequence.
- This “**premature**” termination of trp operon transcription occurs only in the presence of tryptophan-charged tRNA Trp.
- When this premature termination or attenuation occurs, a truncated (140 nucleotides) trp transcript is produced.

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- *The principle of attenuation is that some external event controls the formation of hairpin needed for intrinsic termination*

## Attenuator region:

- It has a nucleotide-pair sequence essentially identical to the *transcription-termination signals* found at the ends of most bacterial operons
- signals contain a G:C-rich palindrome followed by several A:T base pairs.
- Transcription of these termination signals yields a nascent RNA with the potential to form a hydrogen-bonded hairpin structure followed by several uracils

# Regulatory components of the *trpL* region





# 4 regions of Leader sequence:

- (1) nucleotides 60–68
- (2) nucleotides 75–83
- (3) nucleotides 110–121
- (4) nucleotides 126–134

## Alternate secondary structures formed by the trpL mRNA—

either region 1 will pair with region 2 and region 3 with region 4, forming a transcription–termination hairpin

or region 2 will base-pair with region 3, preventing region 3 from pairing with region 4

**The concentration of tryptophan in the cell determines which of these structures will form during the transcription of the trp operon.**

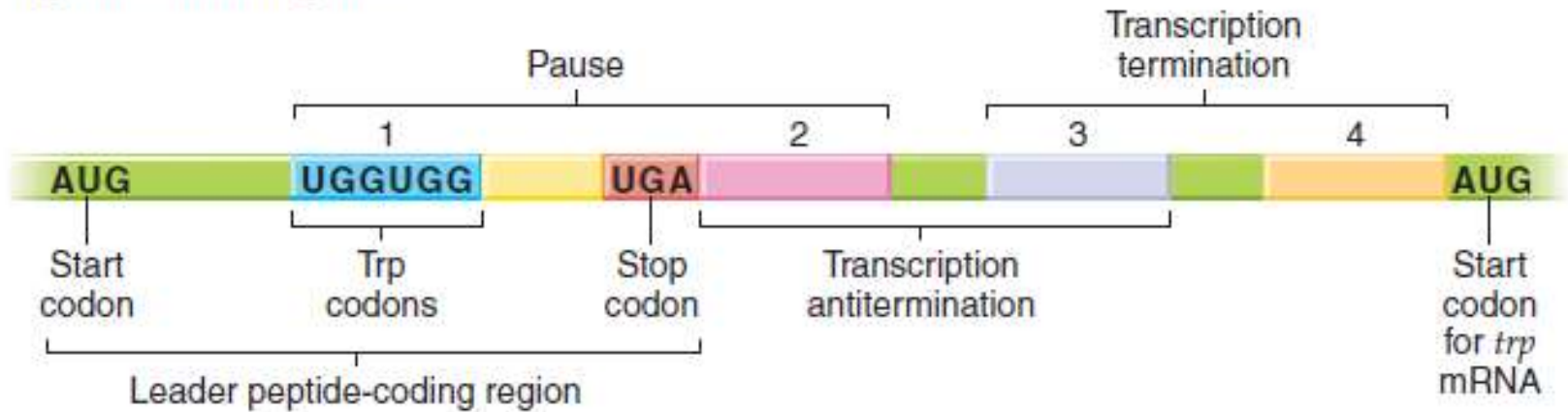
## Crucial three signals :

- The pairing of regions 1 and 2 results in the formation of a **transcription pause signal**
- that of 3 and 4 is a **termination of transcription signal**
- the pairing of 2 and 3 forms an **antitermination signal** for transcription to continue.

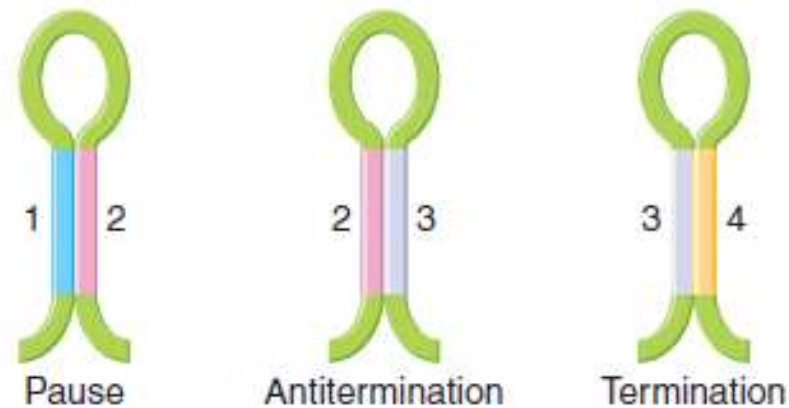


Four regions of the *trp* operon leader mRNA and the alternative secondary structures they can form by complementary base pairing.

Organization of region:

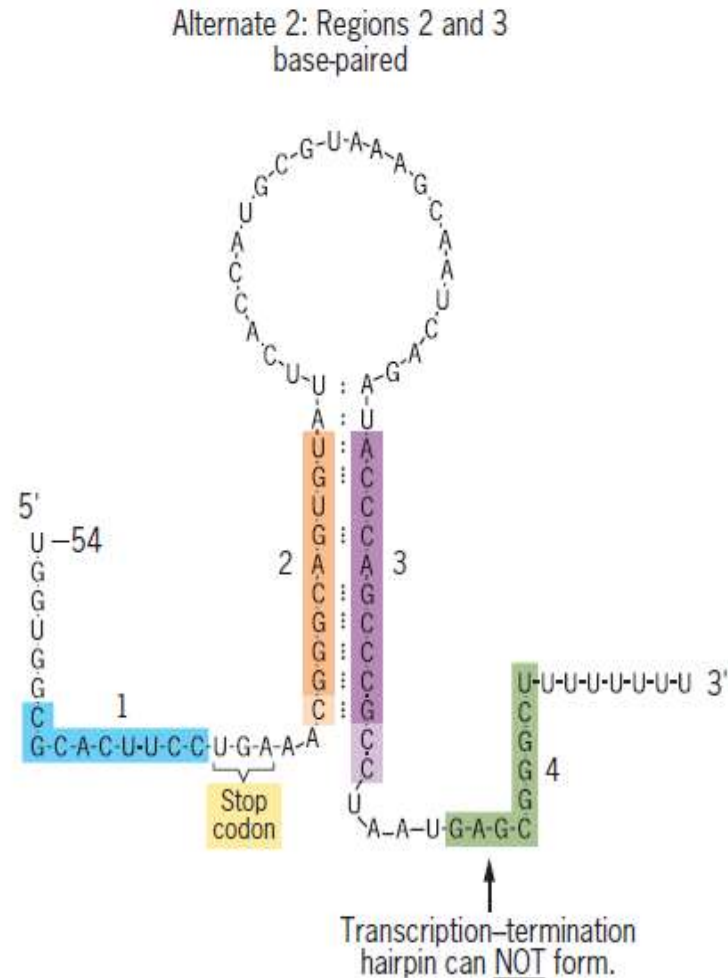
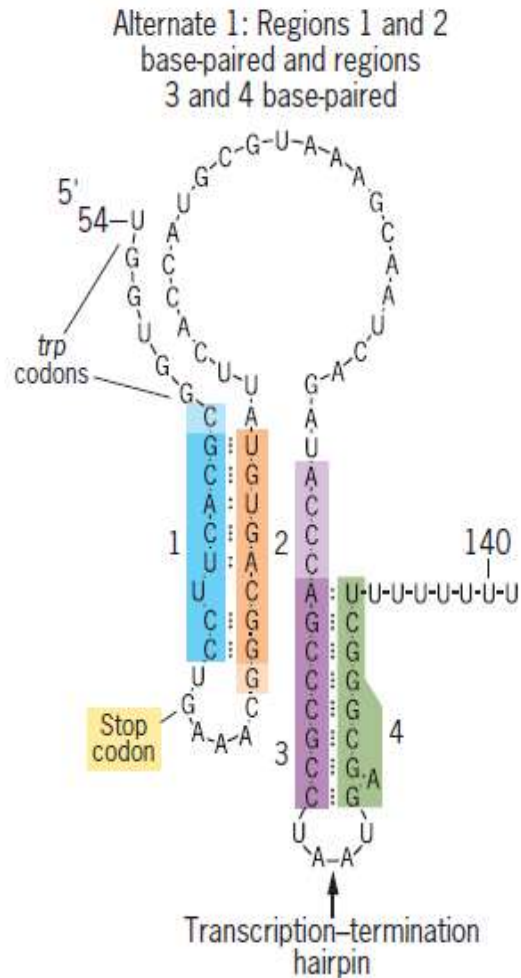


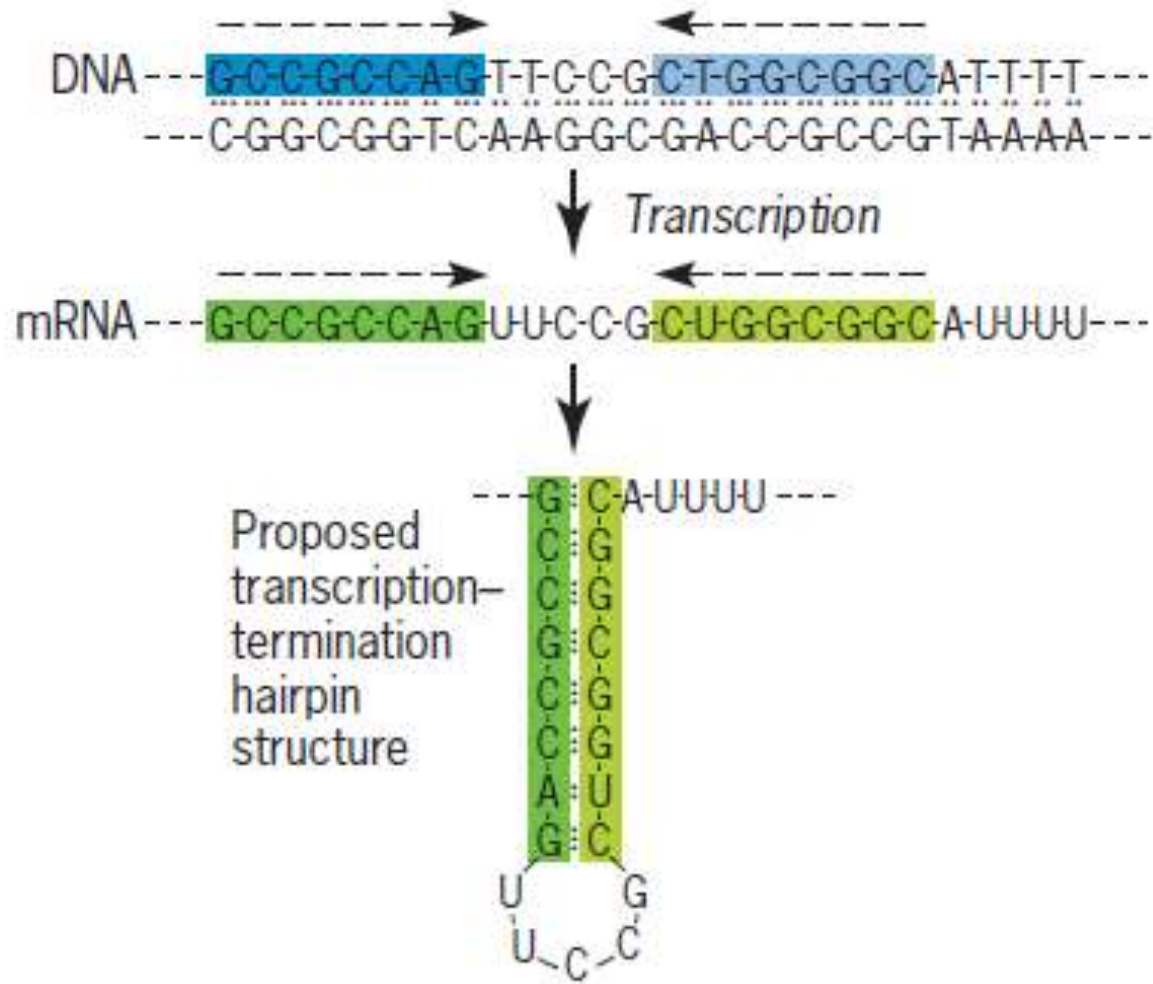
Alternative RNA structures:



# Mechanism :

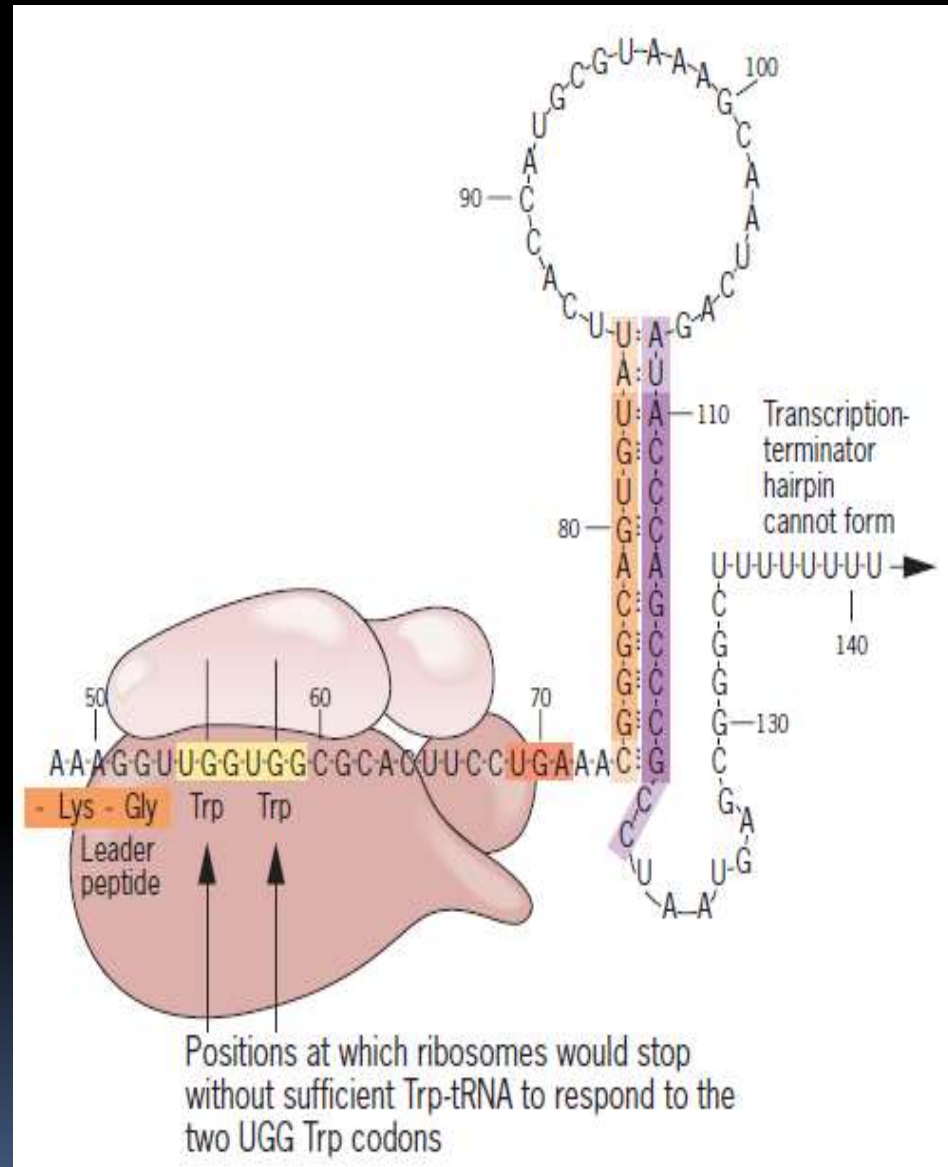
## Alternate secondary structures formed by the *trpL* transcript



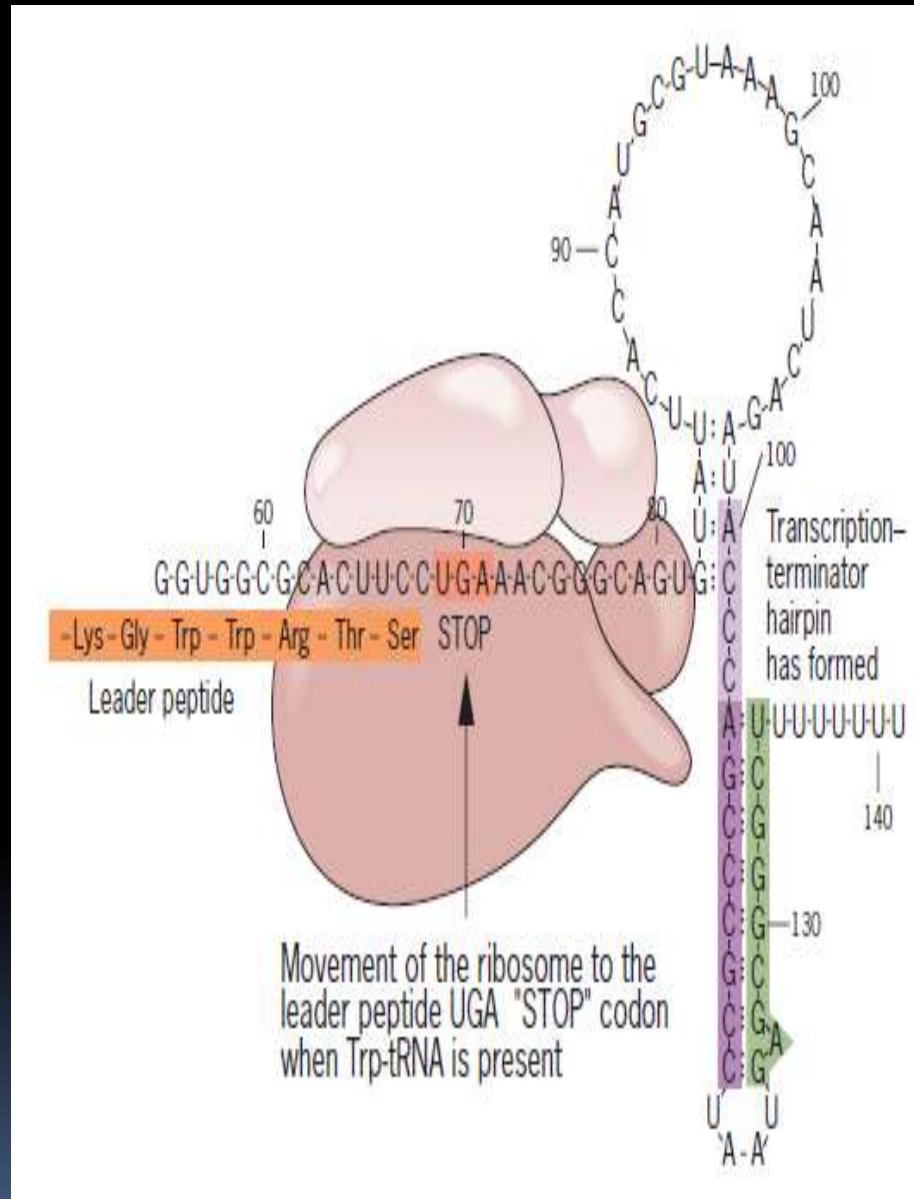


Structure of *trp* operon transcription-termination sequence *t* and formation of the transcription-termination hairpin.

- With low levels of tryptophan, translation of the leader sequence stalls at one of the Trp codons. This stalling allows leader regions 2 and 3 to pair, which prevents region 3 from pairing with region 4 to form the transcription-termination hairpin. Thus transcription proceeds through the entire *trp operon*.



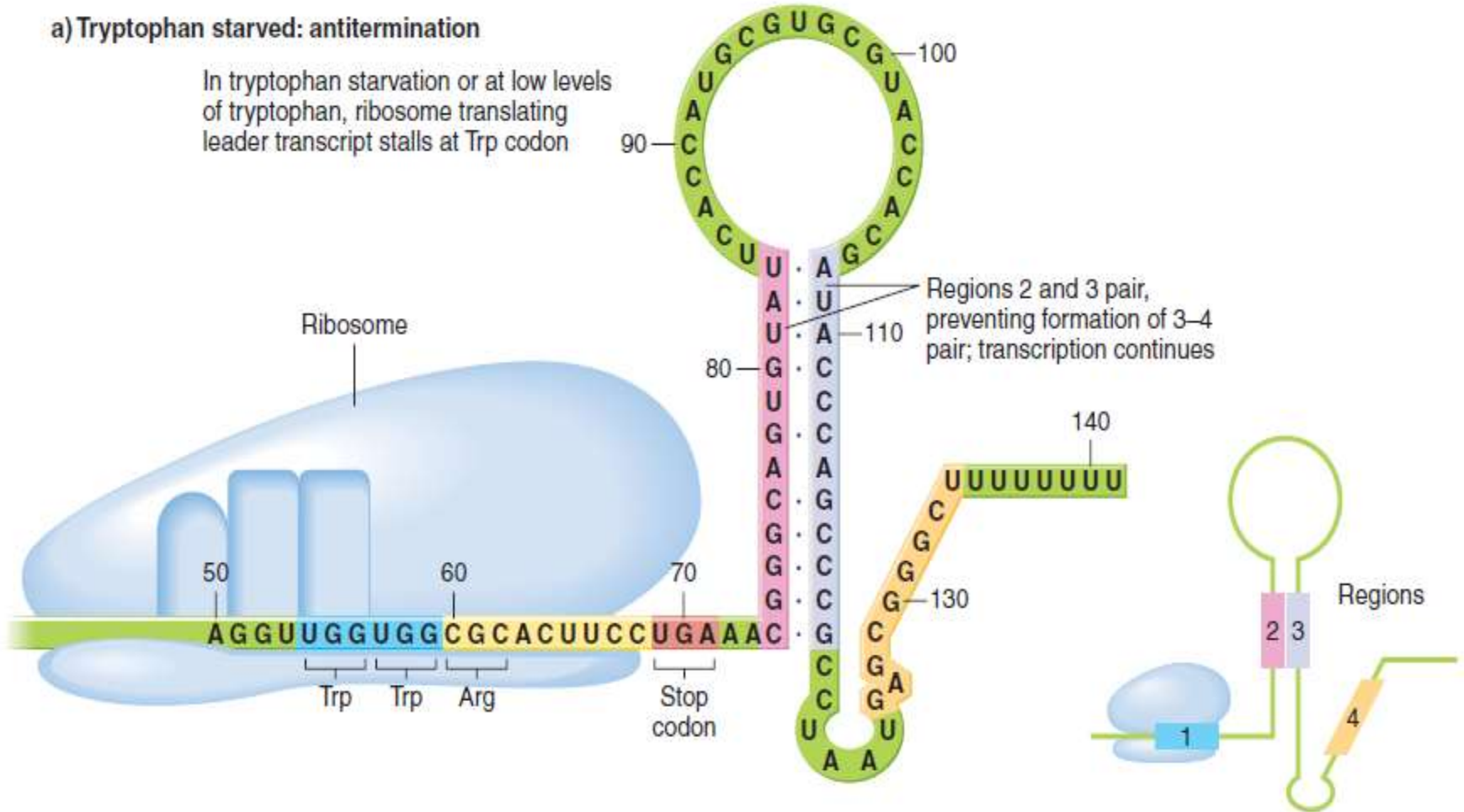
- In the presence of sufficient tryptophan, translation proceeds past the Trp codons to the termination codon and disrupts the base pairing between leader regions 2 and 3. This process leaves region 3 free to pair with region 4 to form the transcription-termination hairpin, which stops transcription at the attenuator sequence.



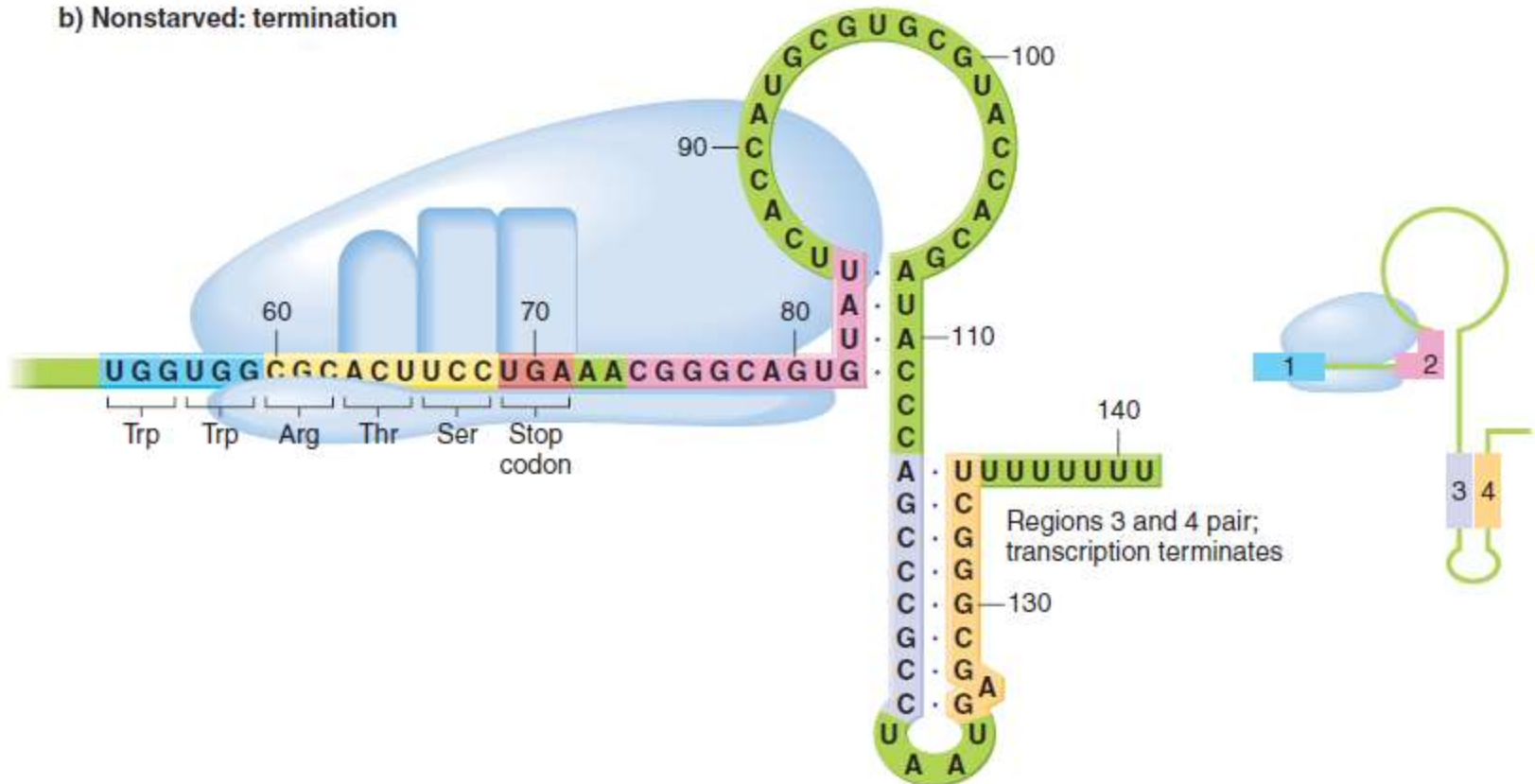
# Model for attenuation in the *trp* operon of *E. coli*.

## a) Tryptophan starved: antitermination

In tryptophan starvation or at low levels of tryptophan, ribosome translating leader transcript stalls at Trp codon

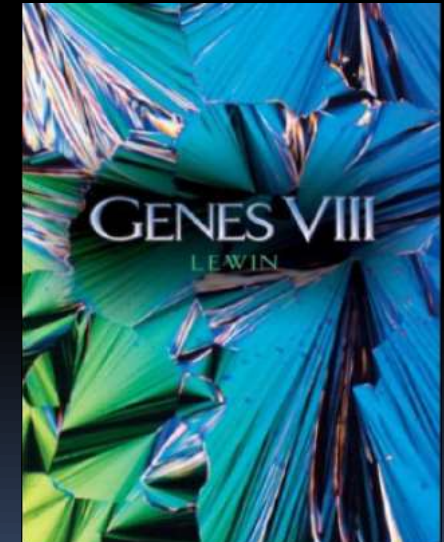
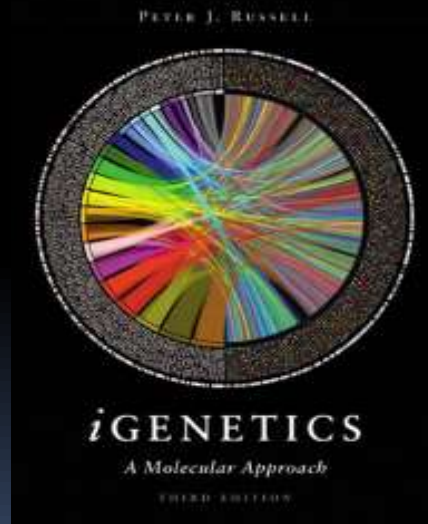
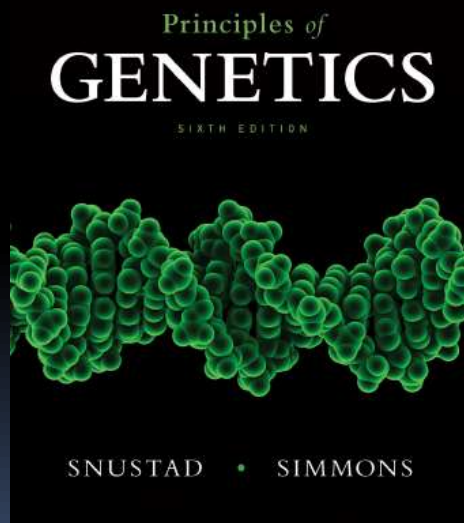


b) Nonstarved: termination



## REFERENCES:

- I. Principles of Genetics by D. Peter Snustad
- II. iGENETICS: A Molecular Approach by Peter J. Russell
- III. Genes VIII by Benjamin Lewin
- IV. Genes IX by Benjamin Lewin







*Thank you for your patience*